### STIC-Biotech/ChemLib

From:

Zeman, Mary

Sent:

Friday, October 25, 2002 8:12 AM STIC-Biotech/ChemLib

To: Subject:

Sequence search 09/970966

Please run a regular search and an oligo search of the following sequences from 09/970966

SEQ ID NO: 199, 210, 211 and 214

please make the minimum window in the oligo searches 10 nucleotides.

please include interference files

Paper printout please.

Thank you, Mary K. Zeman Examiner, 1631 305-7133

CM1 12A17

mailbox: CM1 12D01

199- EFD 09/656,668

**Point of Contact:** Thomas G. Larson, Ph.D. 703-308-7309 CM1, Rm. 6 B 01

Searcher:	arsin
Phone:	
Location:	
Date Picked Up:	11/08/00
Searcher Prep/R	Review:_ <i>40</i>
Clerical:	
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TYPE OF SEARCH:
NA Sequences: 4
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other

VENDOR/COST (where applic.)
STN:
DIALOG:
Questel/Orbit:
DRLink:
Lexis/Nexis:
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AUTHORS TITLE JOURNAL	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	Locus	AX093381	RESULT 1
Xu,J. and Stolk,J.A.  Ovarian tumor sequences and methods of use therefor Patent: WO 0118046-A 199 15-MAR-2001;	Mammalia: Butheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 369)	Homo sapiens	human.		AX093381.1 GI:13509829	AX093381	Sequence 199 from Patent WO0118046.	AX093381 369 bp DNA linear PAT 30-MAK-2001	•	

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                                            CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: National Institutes of Health Intramural
Sequencing Center (NISC),
Gaithersburg, Maryland;
                                                                                                                                                                                     Direct Submission

Direct Submission

Submitted (05-NOV-2001) National Institutes of Health, Mammalian Submitted (05-NOV-2001) Cancer Genomics Office, National Cancer Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                            NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
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AX358762
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This clone was selected for full length sequencing because it passed the following selection criteria: Hexamer frequency ORF analysis, Similarity but not identity to protein.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                            Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zhang, L.-H. and Green, E.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /product="Unknown (protein for MGC:29643)"
/protein_id="AAH17318.1"
/db_xref="GI:16878240"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="MGC:29643 IMAGE:3641660"
/tissue_type="Pancreas, adenocarcinoma"
/clone_lib="NIH_MGC_39"
/lab_host="DH10B-R"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  1524 bp
Sequence 15 from Patent W00208288.
AX362255
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                                                                                                                                                                                                                                                                                                                                      Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7 Kazusa-Kamatari, Kisarazu, Chiba 292-0812, Japan (E-mail:genomics@hri.co.)p, Tel:81-438-52-3975, Fax:81-438-52-3986) NEDO human cDNA sequencing project supported by Ministry of Economy, Trade and Industry of Japan; cDNA full insert sequencing: Research Association for Biotechnology (RAB); cDNA library construction: Helix Research Institute (HRI) (supported by Japan KRy Technology Center etc.); 5'- & 3'-end one pass sequencing: RAB, HRI, and Biotechnology Center, National Institute of Technology and Evaluation; clone selection for full insert sequencing: HRI and RAB; annotation: HRI and RAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AK094501 1832 bp mRNA linear PRI 15-JUL-200: Homo sapiens cDNA FLJ37182 fis, clone BRALZ2001350, weakly similar to Homo sapiens Gz-selective GTPase-activating protein (RGS20)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kawakami,B., Sugiyama,A., Takemoto,M., Sugiyama,T., Irie,R., Otsuki,T., Sato,H., Wakamatsu,A., Ishii,S., Yanamoto,J., Isono Kawai-Hio,Y., Saito,K., Nishikawa,T., Kimura,K., Yanashita,H., Matsuo,K., Nakamura,Y., Sekine,M., Kikuchi,H., Kanda,K., Wagatsuma,M., Muzakawa,K., Kanehori,K., Takahashi-Fujii,A., Oshima,A., Suzuki,Y., Sugano,S., Nagahari,K., Masuho,Y., Nagai
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Mammalia; Eutheria; Primates;
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 /clone="BRALIZ0001350"
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215. .640
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                                                                                                                                                    Location/Qualifiers
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PRI 15-JUL-2002

Isono, Y.,

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SOURCE
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                                                                                                                                                                                                                                                                                                                    Secretory protein or membrane protein Patent: EP 1067182-A 203 10-JAN-2001;
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Sequence 203 from Patent EP1067182.
AX136281
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1. (bases 1 to 1890)
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274. .771
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3 501 c 513 g 407 t
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Score 315;

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                                                                         Local Similarity nes 314; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Millennium Pharmaceuticals, Inc. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cardiovascular and tumorigenic disease using Patent: WO 0181634-A 3 01-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Galvin, K.A. and Rudolph-Owen, L.A. Methods and compositions for the diagnosis and
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VTSALVALPLLFAMGTEYPLVNVPSHAGLTCNRSSTRHHEQPETSNMSICTMLSSRMT
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ARRQTIIFLRLIVVTLAVCWMPNQIRRIMAAAKPKHDWTRSYFRAYMILLPFSETFFY
LSSYINPLLYTVSSQQFRRVFVQVLCCRLSLQHANHEKRLRVHAHSTTDSARFVQRPL
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/db_xref="taxon:9606"
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Pred. No. 7.8e-1
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Sequence 9 from Patent W00118046
AX093191
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The sequence of Homo sapiens
Unpublished (2001)
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MO 63108, USA
4 (bases 1 to 129676)
Waterston, R.H.
                                 Submitted (10-SEP-2000) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St.
                                                                  3 (bases 1 to 129676)
Waterston, R.H.
                                                                                                                                                   Toward a complete human genome Genome Res. 8 (11), 1097-1108 (99063792
                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 129676)
Sulston, J.E. and Waterston, R.
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Submitted (09-JAN-2002) Department of Genetics, Washington
University, 4444 Forest Park Avenue, St. Louis, Missouri 63108,
On Aug 9, 2001 this sequence version replaced gi:14488388.

On Aug 9. Genome Center

Genome Sequencing Center
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        libraries. Genomics 51:1-8. The clone may be obtained either in Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at the Roswell Park Cancer Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Frengen, E., Tatano, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from construction of the content of the co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
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Contact: sapiens@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Center: Washington University Genome Sequencing Center Center code: WUGSC
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1397. .3331
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                 /rpt_family="(TTTTG)n"
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JOURNAL REFERENCE
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                                                                                    2 (bases 1 to 147131)
VanBrunt, A. and Strowmatt, C.
The sequence of Homo sapiens BAC clone RP11-159N20
Unpublished (2001)
3 (bases 1 to 147131)
<u>₩</u> 4
          Submitted (28-SEP-1999) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St. MO 63108, USA
                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 147131)
Sulston, J. E. and Waterston, R.
Toward a complete human genome sequence Res. 8 (11), 1097-1108 (1998)
                                                        Direct Submission
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29414. .29696
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                                                                                                            Submitted (01-MAR-2002) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 63108, USA On Mar 1, 2002 this sequence version replaced gi:13435273.
                                                                                                                                                                                                                                                                 Submitted (23-MAR-2001) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St. Louis,
                                                                                                                                                                                          Direct
                                                                                                                                                                                                              Waterston, R
                                                                                                                                                                                                                                                 MO 63108,
                                                                                                                                                                                                                                                                                                        Direct Submission
                                                                                                                                                                                                                                                                                                                          Waterston, R.H.
                                                                                                                                                                                                                               (bases 1 to 147131)
                 Center project name: H_NH0159N20
                                                                                                                                                                                        Submission
                                                                                                                                                                                                                                                     USA
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NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.

all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality > 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest. This sequence was finished as follows unless otherwise noted:

MAPPING INFORMATION:
Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc Louis

### SOURCE INFORMATION:

donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Frengen, E. Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at http://www.chori.org The RPCI-11 human BAC library was made from the blood of one male pBACe3.6

# NEIGHBORING SEQUENCE INFORMATION:

The clone sequenced to the right is RP11-458A7, 200 bp overlap; clone sequenced to the right is RP11-258B17. Actual start of the clone is at base position 1 of RP11-159N20; actual end is at base position 147131 of RP11-159N20. is at base

clone There are polymorphic base differences RP11-159N20 and RP11-258B17. in the overlap between the

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FEATURES
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                                                 /rpt_family="L1" 984 . .1176
                                                                                      /rpt_family="MIR" 118. .1013
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991. .1176
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/chromosome="2"
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                                                                                                                                                          /clone="RP11-159N20"
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"match to EST
BG184021 (NID:g13705708)"
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7811. .7881
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1525. .1552
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15183. .15321
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1035. .1200
/note="match to 16665. .16728
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9348. .9395
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Submitted (05-JUL-2002) Genome Sequencing Center, 4444 Forest Parkway, St. Louis, MO 63108, USA On Jul 5, 2002 this sequence version replaced gi:21426614.
                                                                        Submitted (14-JUN-2002) Genome Sequencing Center, 4444 Forest Park Parkway, St. Louis, MO 63108, USA 3 (bases 1 to 209885) McPherson,J.D. and Waterston,R.H.
                                                                                                                                               2 (bases 1 to 209885)
McPherson, J.D. and Waterston, R.H.
Direct Submission
                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 209885)
McPherson, J.D. and Waterston, R.H.
The sequence of Mus musculus clone
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AC124493.2 GI:21699722
HTG; HTGS_PHASE1; HTGS_1
                                                                                                                                                                                                                                                                                                                                                                                                                                AC124493 209885 bp DNA line Mus musculus chromosome UNK clone RP23-462P13,
                                                         Direct Submission
                                                                                                                                                                                                       Unpublished
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Db 103249 GTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGAAGGAGCAAAGTGC 103203
                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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Sequencing vector: plasmid; 100%
Sequencing vector: plasmid; 100%
Chemistry: Dye-primer ET; 0% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.990319
Consensus quality: 208484 bases at least Q40
Consensus quality: 208911 bases at least Q30
Consensus quality: 209285 bases at least Q20
Insert size: 206000; agarose-fp
Insert size: 212112; sum-of-contigs
Quality coverage: 11.62 in Q20 bases; sum-of-contigs
Quality coverage: 11.62 in Q20 bases; sum-of-contigs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Web site:http://genome.wustl.edu/gsc/index.shtml
Contact: submissions@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Center: Washington University Genome Sequencing Center Center code: WUGSC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               * NOTE: This is a 'working draft' sequence. It currently consists of 5 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence.
Mus musculus brain cDNA, clone MNCb-0671.
AB041649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  as soon as it is available and the accession number will be preserved.
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11432
28618
28718
49455
49555
107385
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                                                                                                                                                                                                                                                                                                                                                                                                                                49555. .107384 // note="assembly_name:Contig11" 107485. .209885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="assembly_name:Contig9"
28718. .49454
/note="assembly_name:Contig10"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1. .209885
                                                                                                                                                                                                                                                                                                                                                                                 /note="assembly_name:Contig12"
44784 c 46164 g 58745 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:10090"
/chromosome="UNK"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="assembly_name:Contig8"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="RP23-462P13"
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11431: gap of unknown length
28617: contig of 17186 bp in length
28717: gap of unknown length
49454: contig of 20737 bp in length
49554: gap of unknown length
107384: contig of 57830 bp in length
107484: gap of unknown length
209885: contig of 102401 bp in length.
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100.0%;
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                                                                                                                                                                                                                                                                                            Score 47;
Pred. No.
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                                                                                                                                                                                                                                                                                         DB 2; L
. 1.9e-16;
                                                                                                                                                                                                                                                                                                                Length 209885;
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                                          ACCESSION
                                                                                    DEFINITION
                                                                                                                               RESULT 14
AL512284/c
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                 VERSION
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                                                                                                                                                                                                                             394 CAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGA 431
                                                                                                                                                                                                                                                                        50 CAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGA 87
                                                                                                                                                                                                                                                                                                                                                Local Similarity
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fis (full insert sequence).
Mus musculus (strain:C57BL) adult female cDNA to mRNA,
clone_lib:Sugano mouse brain mncb clone:MNCb-0671.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ist strand cDNA was primed with an oligo(dT) primer lst strand cDNA was primed with an oligo(dT) primer larger stranded cDNA was synthesized using specific 5′ and 3′ primers and amplified by PCR. The PCR product was digested with SfiI and size selection was performed to exclude fragments <1.5kb. The SfiI-digested PCR product was cloned into distinct DraIII sites of pwEl8s-FL3. XhoI sites just outside the DraIII sites can be used to isolate the cDNA insert. Library was constructed by Sugano et al. (University of Tokyo, Institute of Medical Science). Custom primer used for sequencing (5′ end primer [CTGCTCTARAAGCTGCG]; 3′ end primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (13-APR-2000) Katsuyuki Hashimoto, National Institute of Infectious Diseases, Division of Genetic Resources; 23-1, Toyama 1-chome, Shinjuku-ku, Tokyo 162-8640, Japan (E-mail:khashi@nih.go.jp, URL:http://www.nih.go.jp/yoken/genebank/.rel:81-3-5285-1111(ex.2120), Fax:81-3-5285-1181)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Osada,N., Kusuda,J., Tanuma,R., Ito,A., Hirata,M., Sugano,S. and Hashimoto,K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                            38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      made by oligo-capping method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               isolation of full-length cDNA clones from mouse brain cDNA library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A part of this sequence is reported in AU035640 
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hashimoto,K., Osada,N., Kusuda,J. and Sugano,S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vector:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Direct Submission
                                                                                  Human DNA sequence from clone RP11-248M22 on chromosome 10
                                       complete sequence
AL512284
                 AL512284.20 GI:18477310
                                                                                                                AL512284
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lame: Sugano mouse brain mncb
                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                          /protein_id="BAA95101.1"
/protein_id="BAA95101.1"
/db_xref="GI:7670500"
/translation="MMVLGIAATFCGLFWLPGLALQIQCYQCEEFQLNNDCSSPEFIV
NCTVNVQDMCQKEVTEQSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRPKKRGSSASAIRPGLLTTLLFFHLALCLAHC"
a 487 c 511 g 408 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Mus musculus"
/strain="C57BL"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="Sugano mouse brain mncb"
/dev_stage="adult"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note-"unnamed protein product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 clone="MNCb-0671"
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FEATURES

BASE COUNT ORIGIN

Matches

RESULT 13 AB041649

PRI 31-JAN-2002

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ACCESSION DEFINITION

SOURCE

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                                                                                                                                                                                                                                                                                                                                                                     38431 CAGTCCTTCTGCTCCCAGGGA 38410
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                        *** SEQUENCING IN PROGRESS *** SEQUENCING IN PROGRESS ***.
AC118794
                                                                                                                                              AC118794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        corresponding to the overlapping clone, as we submit sequences with only a small overlap as described above.

This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least assembly was confirmed by restriction digest. The following abbreviations are used to associate primary accession numbers given in the feature table with their source databases: Em: EMBL; Sw:, survespanner. The manufacture of the survespanner.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IMPORTANT: This sequence is not the entire insert of clone RP11-248M22 It may be shorter because we sequence overlapping sections only once, except for a short overlap. The true left end of clone RP11-248M22 is at 1 in this sequence. The true left end of clone RP11-495F22 is at 11921 in this sequence. The true right end of clone RP11-481F7 is at 53678 in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    http://www.sanger.ac.uk/Projects/C_elegans/wormpep This sequence was generated from part of bacterial clone contigs of human chromosome 10, constructed by the Sanger Centre Chromosome 10 Mapping Group. Further information can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                http://www.sanger.ac.uk/HGP/Chr10
RP11-248M22 is from the library RPCI-11.1 constructed by the group
of Pieter de Jong. For further details see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SWISSPROT; Tr:, TREMBL; Wp:, WORMPEP; Information on the WORMPEP database can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Where differences are found these are annotated as variations together with a note of the overlapping clone name. Note that the variation annotation may not be found in the sequence submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk On Feb 1, 2002 this sequence version replaced gi:18476636. During sequence assembly data is compared from overlapping clones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="RP11-248M22"
/clone_lib="RPCI-11.1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Carroy, C., Carter, Burkell, Burgal, N.C., Carrol, T.F., Carter, M., Cavacos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Coyle, M.D., Dathorne, S.R., David, R., David, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., David, R., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Flagy, N., Ford, J., Foster, P., Frantz, P., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gorrell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K., Harris, C., Harris, K., Hart, M., Haylak, P., Hale, S., Hamilton, K., Harris, C., Harris, C., Harris, K., Hart, M., Holloway, C., Hollins, B., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C., Kureshi, A., Landry, N., Leal, B., Lewis, L., Li, J., Li, Chitarge, O., Lieu, C., Liu, J., Liu, W., Loulseged, H., Maheshwari, M., Mapua, P., Martin, R., Martindale, A., Martinez, E., Maheshwari, M., Mapua, P., Martin, R., Martindale, A., Martinez, E., Maher, G., Miner, S., Michell, T., Mohabbat, K., Morgan, M., Morris, S., Massey, E., Mawhiney, E., McLeod, M.P., Meador, M., Nguyen, A., Nguyen, N., Nguyen, N., Newtson, J., Newtson, J., Newtson, M., Savery, G., Scher, S., Sonaike, T., Sonaike, T., Sonaike, T., Savery, G., Stenley, H., Stone, H., Wang, S., Ward-Moore, S., Warren, R., Washington, C., Watlington, S., Wulliamson, A., Warren, R., Washington, C., Watlington, S., Warren, R., Washington, S., Warren, R., Washington, S., Warren, S., Shan, R., Wang, O., Watlington, S., Warren, R
                                                                                                                                                                                                                                                                                                                                                                                        Submitted (18-JUL-2002) Human Genome Sequencing of Molecular and Human Genetics, Baylor College Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                    On Jul 14, 2002 this sequence version replaced g1:20387310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (21-APR-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
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Buhay, C., Burch, P., Burkett, C., Burrell, K.L., Byrd, N.
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                                                Center project Information Center project name: GVHE Center clone name: CH230-179C24
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                                                                                                                                                            Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
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NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 69 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence.
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Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Chemistry: Dye-terminator Big Dye; 90% of reads
Chemistry: Dye-primer Big Dye; 10% of reads
Chemistry: Dye-primer Big Dye; 10% of reads
Assembly program: Phrap; version 0.990399
Note:data AC066614 [Drafting center UWMSC] and AC066611
[Drafting center UWMSC] were added for finishing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
Submitted (10-UUL-2001) Multimegabase Sequencing Center, Institution Systems Biology, 4225 Roosevelt Way NE, Suite 200, Seattle,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (22-FEB-2000) Multimegabase Sequencing Center, University of Washington, PO BOX 357730, Seattle, WA 98195, USA 3 (bases 1 to 157042)
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On Jul 10, 2001 this sequence version replaced gi:13162487
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: leerowen@systemsbiology.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Web site: http://chroma.mbt.washington.edu/msg_www
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ----- Summary Statistics
                                                                                         Conservative
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                                                                                                                                                                                 143570. 157042
/note="overlap with RP11-348P16 AC066611.
a 28493 c 28053 g 47690 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /Clone_lib-"RPCI Human BAC library 11"
/note-"This clone overlaps RP11-8068 14 and
/note-"This clone overlaps RP11-8068 14.
RP11-348P16 AC066611. Data from overlapping BACs were
combined and the consensus sequence determined from
RP11-232J12 to the extent possible."
                                                                                                                                                                                                                                                       /note="low quality data."
143570. .157042
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124490. .124492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note="overlap with RP11-80G8 AC066614"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 to 157042)
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                                                                                                             100.0%;
                                                                                                             5.4%; Score 20; DB 9; 100.0%; Pred. No. 3.9;
                                                                                                                                                                                                                                                                                                                                                                                                        .4280C
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                                                                                         0
                                                                                         Mismatches
                                                                                       0,
                                                                                                                                 Length 157042;
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                                                                                    0;
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FEATURES

Location/Qualifiers

Exon/Intron boundaries of identified genes were chosen if there were canonical splice junctions that maintained sequence continuity across the splice junctions.

The repeat regions shown were identified using RepeatMasker by Adrian Smit.

Sequence similarities were identified using Powerblast by Jinghui

source

COMMENT

REFERENCE

JOURNAL TITLE JOURNAL

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (01-MAR-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Dec 1, 2000 this sequence version replaced gi:11321701. Sequencing is completed to a minimum standard of double strand coverage with a minimum of 2 clones and 2 reads with no ambiguities or 2 chemistries with a minimum of 2 clones and 3 reads with no ambiguities. If the sequence quality does not meet this standard, it will be indicated in the annotation.
                                                                                                                                                                                                                                                                                                             Submitted (05-DEC-2000) Human Genome Sequencing of Molecular and Human Genetics, Baylor College Baylor Plaza, Houston, TX 77030, USA (bases 1 to 171146)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fernandez,C., Ferraguto,D., Forcum Tansey,J., Gill,R., Gorrell,J.H., Gunaratne,P., Haller,G., Hernandez,J., Hogues,M., Hosak,H., Hubur,J., Jackson,L., Jia,Y., Kelly,J., Kelly,S., Hosak,H., Hou,X., Huber,J., Jackson,L., Jia,Y., Kelly,J., Kelly,S., Kovar,C., Liu,J., Liu,W., Loulseged,H., Lozado,R.J., Martin,R., Massey,E., McLeod,M.P., Mei,G., Moore,S., Morgan,M., Morrits,S., Massey,E., McLeod,N.P., Mei,G., Moore,S., Morgan,M., Morrits,S., Nelson,A., Nguyen,R., Nguyen,N., Oguh,M., Parish,B., Perez,L., Reiter,D., Say,J., Shen,H., Vasquez,L., Watlington,S., Williamson,A., Wrensford,G., Zhou,X., Bouck,J., Hodgson,A., Muzny,D.M., Rives,M., Scherer,S., Sodergren,E., Weinstock,G., Muzny, D.M., Rives,M., Scherer,S., Sodergren,E., Weinstock,G., Muzny,K., and Gibbs,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of Molecular and Human
Baylor Plaza, Houston,
4 (bases 1 to 171146)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (10-MAY-2000) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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Direct Submission
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Direct Submission
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Direct Submission
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                                                                                                                                                                                                                                                                 Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TX 77030,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
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Brooks,A., Brown,J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Center, Department
of Medicine, One
                                                                                                                                                                                                                                                                                                                                                                            Center, Department of Medicine, One
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complement(9725. .9861)
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4033. .4231
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                                                                                                                                                                                                                                                                       rpt_family-"B1_MM"
                                                                                                                                                                                                                                                                                                  rpt_family="A-rich"
                                                                                                                                                                                                                                                                                                                                rpt_family="B3"
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                                   _family="CT-rich"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 _family="(CGG)n"
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                                                                                            _family="CT-rich"
                       .13842
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                             DEFINITION
ACCESSION
                                                          AC099049/c
                                                                                                                                 Db 156695 CTCTGCCGGGTACCAGTCCT 156714
                VERSION
                                                                                         RESULT 18
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Best Local :
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                                                                                                                                                                                                    Local
180339 bp DNA linear PRI 12-JAN-
Homo sapiens chromosome 3 clone RP11-291P10, complete sequence.
AC099049 AC021722
AC099049.2 GI:18139443
                                                                                                                                                                                          l Similarity
20; Conserv
                                                                                                                                                                                       5.4%; Score 20; DB 10; Length 171146; ilarity 100.0%; Pred. No. 3.9; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                             /rpt_family="c-rich"

complement(26401 . 26552)

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complement(26725 . 26915)

/rpt_family="83A"

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20895
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          _family="AT_rich"
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0. .21087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      _family="(CA)n"
                                                                                                                                                                                                                                                                                                                                                                                 _family="B1-F"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                _family="PB1D10"
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. 9608

				COMMENT	TITLE JOURNAL	REFERENCE AUTHORS	JOURNAL REFERENCE AUTHORS TITLE JOURNAL	REFERENCE AUTHORS	KEYWORDS SOURCE ORGANISM
This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., Phred quality — 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one MI3 subclone; and the assembly was confirmed by restriction digest.  Sequence Validation: This sequence has been validated by Multiple Complete Digest fingerprinting. Comparison of the experimentally derived digest fragments with sequence-predicted fragments is given below. The electronically-digested sequence consists of both insert and vector, in order to accurately represent the entire circular BAC. Small fragments below a variable cutoff (approximately 400-800 bp) are not resolved in the fingerprint and hence do not appear	Sequence Quality Assessment: This entry has been annotated with sequence quality estimates computed by the Phrap assembly program. All manually edited bases have been reduced to quality zero. Quality levels above 40 are expected to have less than 1 error in 10,000 bp. Base-by-base quality values are not generally visible from the GenBank flat file format but are available as part of this entry's ASN.1 file.	rlappin	Code: UWGC  e: http://www.genome.washington.ed  uwgchtgseu.washington.edu  g Center: BCM  Project Information  project name: chr-3  clone name: RP11-291P10 (bc0345)  ing vector: unknown; 60% of reads  fing vector: plasmid; 40% of reads  ry: Dye-terminator Big Dye; 11%  ry: Dye-terminator Big Dye; 11%  gy: Dye-terminato	On Jan 12, 2002 this sequence version replaced 91:16/9900/ Genome Center Center University of Washington Genome Center	vers	339) .V., Zhou,Y., Karen,P.K.A.	Unpublished 2 (bases 1 to 180339) 2 (bases 1 to 180339) Kaul,R.K., Olson,M.V., Raymond,C. and Haugen,E.D. Direct Submission Submitted (08-NOV-2001) Genome Center, University of Washington, DAY 353145 (Bentle Wa 98195 USA	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 180339)  1 (bases 1 to 180339)  Kaul, R.K., Olson, M.V., Zhou, Y., James, R.A., Rouse, G., Wu, Z., Saenphimmachak, C., Karen, P.K.A., Raymond, C. and Haugen, E.D.  Direct Submission	

in the table. There are no significant remaining discrepancies between the experimental and predicted values. Uniquely ordered fragments are separated by dashed lines.

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AUTHORS

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190914 ns chromosome 3 clo ed pieces. 4 GI:22203847 pHASEI; HTGS_DRAFT. ns Metazoa; Chordata; Eutheria; Primates; 1 to 190914)		5.4%; 100.0%; 7ative	6536	13421	1373	<800	<800	2932	963	2314	5278	2932	2932	4708	8247	3499	2606	4986	3247	800	5278	3719	4422	<800	843	1230	843
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HTG 24-AUG-2002 DRAFT SEQUENCE, Euteleostomi;		Gaps 0;	981	<800	4606	<800	<800	1231	4606	5042	878	2052	3053	2052	2673	3257	2052	2052	5042	7809	1160	6296	4154	3257	5042	1347	1573
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Bunay, C., Butch, F., Butch, C., Butlett, C., Carron, T.F., Carter, M., Cavacos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davil, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., Davil, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., Davil, R., L., Davis, C., Davy-Carroll, L., Dederich, D.A., Davil, R., L., Ding, Y., Dinh, H.H., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delaney, K.R., L., Delaney, K., Bord, J., Boster, P., Frantz, P., Falls, T., Ferraguto, D., Flagg, N., Ford, J., Boster, P., Frantz, P., Falls, T., Ferraguto, D., Flagg, N., Ford, J., Boster, P., Frantz, P., Harris, C., Edwards, C.C., Elhaj, C., Escotto, M., Earlhart, C., Edwards, C.C., Elhaj, C., Escotto, M., Earlhart, C., Edwards, C., Elhaj, C., Escotto, M., Frantz, P., Hawes, A., Hernandez, J., Hogues, M., Holloway, C., Hollins, B., Hernandez, O., Hodgson, A., Hogues, M., Holloway, C., Hollins, B., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Jacobson, B., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C., Karlsson, E., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C., Karlsson, E., Lichtarge, O., Lieu, C., Liu, J., Liu, W., Loulseged, H., Li, J., Liu, W., Loulseged, H., Li, J., Liu, W., Loulseged, H., Li, J., Liu, W., Loulseged, H., Massey, E., Machaley, N., Meador, M., Martinez, E., Michell, T., Mohabbat, K., Morgan, M., Morris, S., Miner, G., Miner, Z., Mitchell, T., Mohabbat, K., Morgan, M., Morris, S., Mogan, N., Noylen, A., Nguyen, A., N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (04-MAR-2000) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 190914)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (24-AUG-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Aug 11, 2002 this sequence version replaced g1:9438348.
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Direct Submission
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Direct Submission
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Center project name: HAJK
Center clone name: HAJK
Center clone name: RP11-442C9
Center clone name: RP11-442C9
Center clone name: RP11-442C9
Center clone name: MP11-442C9
Chemistry evector: M13;
Sequencing vector: M13;
Consensus quality: 16241 bases at least Q30
Consensus quality: 181136 bases at least Q20
Estimated insert size: 191206; sum-of-contigs estimation
Quality coverage: 3x in Q20 bases; sum-of-contigs estimation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Web site: http://www.hgsc.bcm.tmc.edu/
Drafting Center Code: BCM
Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Center: Baylor College of Medicine
Center code: BCM
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JOURNAL REFERENCE

TITLE

TITLE AUTHORS

JOURNAL

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COMMENT

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                                  Query Match
             Matches
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                        Local Similarity
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NOTE: This is a 'working draft' sequence. It currently consists of 25 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one M13 subclone; and the assembly was confirmed by restriction digest. The following abbreviations are used to associate primary accession numbers given in the feature table with their source databases: Em:, EMBL; Sw:, SMISSPROT; Tr:, TREMBL; Wp:, WORMPEP; Information on the WORMPEP database can be found at
                                                                                                                                                                                                                                                                                                                                                         20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (04-APR-2002) Wellcome Trust Sanger Institute, Hinxton, Cambridgeshire, CB10 15A, UK. E-mail enquiries: humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk On Apr 7, 2002 this sequence version replaced gi:15591682. During sequence assembly data is compared from overlapping clones. Where differences are found these are annotated as variations together with a note of the overlapping clone name. Note that the variation annotation may not be found in the sequence submission corresponding to the overlapping clone, as we submit sequences with may a small overlap as described above.
                                               AX392969.1
                                                                                                                          AX392969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fttp://www.sanger.ac.uk/Projects/C_elegans/wormpep RP23-180B18 is
from the RPCI-23 Mouse PAC Library
constructed by the group of Pieter de Jong.
For further details see http://www.chori.org/bacpac/home.htm
human
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/clone_lib="RPCI-23"
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/db_xref="taxon:10090"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 13 from Patent W00214358.
AX392971
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HLRSLGALRLRHLAIASLEDQNFRCLPGLHLEIDNWPLLEEVAAGSLRGLKLTSLSV 
THTNITAVPAAALRHQAHLTCLNLSHNPISTVPRGSFRDLVRLRELHLAGALLAVVEP 
QAFLGLRQIRLLNLSNNLLSTLEESTFHSVNTLETLRVDGNPLACDCRLLWIVQRRKT
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LCRAEGEPAPTVAWVTPQHRPVTATSAGRARVLPGGTLEIQDARPQDSGTYTCVASNA
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49. .1827
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              RLRVLRLRGNQLKLIPPGVFTRLDNLTLLDLSENKLVILLDYTFQDLHSLRRLEVGDN
DLVFVSRRAFAGLLALEELTLERCNLTALSGESLGHLRSLGALRLRHLAIASLEDQNF
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/db_xref="G1:19701019"
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/translation="wyrchicytispilllepanpppaggcparcectvotravactrrr
LTAVPDGIPAETRILELSRNRIRCLNPGDLAALPALEELDLSENAIAHYEPGAFANLY
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RRLPGLLHLEIDNWPLLEEVAAGSLRGLNLTSLSVTHTNITAVPAAALRHQAHLTCLN
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Sequence 9 from Patent WO0248361.
AX463719
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                                                                                l Similarity
19; Conser
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                                                                                                                                                                                                                                                 ATSAGRARVLÞGGTLEIQDARÞQDSGTYTCVASNAGGNDTYFATLTVRÞEÞAANRTÞG
EAHNETLAALRAÞLDLTTILVSTAMGCITFLGVVLFCFVLLFVMSRGRGQHKNNFSVE
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Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
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                                                                                                                                                                                                                           YSFRKVDGPAAAAGQGGARKFNMKMI"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /codon_start=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="unnamed protein product"
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Primates;
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AF213399

2349 bp

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PLN 10-DEC-2000

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D49515/c
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                                                                                                                                                                                                                                                                                                                                                    304 GCCAAGGCCCAAGAAAAGG 322
                                                                                                                                              complete cds.
D49515
D49515.1 GI:1902918
Kawasaki, K., Kubo, T. and Natori, S.
                                              Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Neoptera; Orthopteroidea; Dictyoptera; Blattaria; J
                                                                                      Periplaneta americana fat body cDNA to mRNA, clone:PL25. Periplaneta americana
                                                                                                                                                                                                    American cockroach clone
                                                                                                                                                                                                                           D49515
                                  Blattidae; Periplaneta.
                                                                                                                         lectin-related protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (07-DEC-1999) Horticulture and Landscape Architecture, Purdue University, 1165 Horticulture Building, West Lafayette, I
                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tobacco and Arabidopsis SLT1 suppresses salt sensitivity of calcineurin-deficient yeast mutants through the regulation of ion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 2349) Matsumoto, T.K., Pardo, J.M., Takeda, S., Bressan, R.A. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Asteridae; euasterids I; Solanales; Solanaceae; Nicotiana.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished
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                       (sites)
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PIKCKKLLNGSGLNLSTQPSNHSNGDAMDLLLPVNGKRKKDFFSNGNEEECYLDVNPP
TSYQLEDLEIHPNEFYWLNEFSGVMRDVYGPVTAAKSIYEDDEGYLIIISLPFVDLQR
VKVSWRNTLTHGIIKVSCLSTSRISFIKRQNRTFKLEASSSEHCPAGEFVREIPLSTR
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SIWGANFFFSFYFKPVLNEKSKPKMVRDSNGVSGFDKSDLQLDVFMVQHDMENMYMWV
FKDRPENALGKNQLRSYMNGHSRQGERPFFSADRGFIRSHRNQRKHYRGLSNPQCVH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product="SLT1 protein"
/protein_id="AAG39002 1"
/db_xref="GI:11611667"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /translation="MVDTLLKTLSMENNHHPSTLLSMDSSASSSHDELDLEMNRQVVI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Nicotiana tabacum"
/cultivar="W38"
/db_xref="taxon:4097"
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e PL25 mF
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                                                  Blattaria; Blattoidea;
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                                                                                          Submitted (25-AUG-1998) Kenzo Sato, Tottori University Faculty of Medicine, Department of Molecular Biology; 86 Nishimachi, Yonago, Tottori 683-8503, Japan (E-mail:kensato@grape.med.tottori-u.ac.jp. Tel:+81-859-34-8039, Fax:+81-859-34-8274)
                                                                                                                                                                                                        Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                Saito, K., Nakamura, T., Komoda, H., Hori, N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AB017043.1 GI:3461893
HNF-3G; hepatocyte nuclear factor
                                                                                                                                                                                                                                                   Saito, K., Nakamura, T., Komoda, H., Hori, N., Adachi, K.,
                                                                                                                                                                                                                                                                                                        Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rattus norvegicus
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                                                                                                                                                                                                                                                                                                                                factor-3gamma (HNF-3G)
                                                                                                                                                                                                                                                                                                                                                   Isolation and characterization of the rat hepatocyte nuclear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kawasaki,K., Kubo,T. and Natori,S.

Presence of a lectin family in Periplaneta americana, revealed by analysis of cDNA clones of lectins related to Periplaneta lectin
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/organism="Rattus norvegicus"
/strain="Wistar"
                                                                        Location/Qualifiers
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1 512 c 561 g 823 t 1 others
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PLHVNVDHTAVLCERSDSVYIVATISAPPRSPSAGYELLPGLGYYKFHTDYKNWYDAR
KTCIQEGAHLAVINSETESKALLKLWLPHPKKNENDWRNDWAHIGFHDHYTEGQFVTIF
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/product="lection-related protein"
/protein_id="BAA18916.1"
/db_xref="GI:1902919"
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/db_xref="taxon:6978"
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3 (bases 1 to 22752)
DOE Joint Genome Institute and Stanford Human Genome Center Direct Submission
                                                                                                                                                               Quality: Phrap Quality >=40 100% of Sequence; Estimated Total Number of Errors is 0.

NOTE: This insert is not the entire sequence of the clone. It is clipped at the overlaps with AC004152 and AC005258. The number of bases overlapped with AC004152 is 7297 and with AC005258 is 10523.
                                                                                                                                                                                                                                                                                                      Submitted (01-MAR-2002) DOE Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA On Mar 1, 2002 this sequence version replaced gi:17646901. Draft Sequence Produced by DOE Joint Genome Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 (bases 1 to 22752)
DOE Joint Genome Institute and Stanford Human Genome Center.
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DOE Joint Genome Institute and Stanford Human Genome Center.
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                                                                                                                                                                                                                                                 Finishing Completed at Stanford Human Genome Center www.shgc.stanford.edu
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2996. .3914
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2695. .2699
2769. .2774
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/protein_id="BA332534.1"
/db_xref="G1:4432776"
/translation="WASGIPGLYASGLNSGRVPVERPKRSVPPGGPQPGRVGAQARGM
/translation="WASGIPGLYASGLNSGRVPVERPKRSVPPGGPQPGRVGAQARGM
LGSYKMEAHDLAEWSYYPEAGE"
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<2798. .>2995
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                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="19"
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Burkhart-Schultz, K., Gordon, L., Kyle, A., Ramirez, M., Stilwagen, S.,
Bhan, H., Velasco, N., Garnes, J., Danganan, L., Poundstone, P.,
Christensen, M., Georgescu, A., Avila, J., Liu, S., Attix, C.,
Andreise, T., Trankheim, M., Amico-Keller, G., Coefield, J., Duarte, S.,
Lucas, S., Bruce, R., Thomas, P., Quan, G., Kronmiller, B., Arellano, A.,
Montgomery, M., Ow, D., Nolan, M., Trong, S., Kobayashi, A., Olsen, A.O.
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AC004152.1 GI:2894631
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Direct Submission
Submitted (18-FEB-1998) Joint Genome Institute, Lawrence Livermore Submitted (18-FEB-1998) Joint Genome Institute, CA 94551, USA National Laboratory, 7000 East Ave., Livermore, CA 94551, USA Map and sequence oriented from p telomere to centromere.
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                                                                                                                                                                                                                                                                             /note="predicted exon, profiame: 1, quality: good, s complement(150. 227) /rpt_family="MIR" complement/1100
                                                                                                                                                              /rpt_family="AluSq/x"
complement(1845. .2140)
/rpt_family="AluX"
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                                                                                                                                                                                                                                                         complement(1189. .1234)
/rpt_family="MIR"
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/clone="fos37308"
complement(3551. .3848)
                                          /rpt_family="AluSx" 2993. .3293
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                         /rpt_family="AluY"
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/rpt_family="Alusc"
complement(12479. .
                                                                                                                                                                                                               complement(10491..10608)
/rpt_family="AluJ"
complement(10667..10921)
                                                                                                                                 DNA for antizyme, complete cds"
                                                                                                                                            /note="BLASTN similarity to D89870 (1. .10524); match:
0.99, score: 2.1e-209; database searched: nt; Homo sap
                               /rpt_family-"MER20"
12211. .12270
                                                                                                                                                                                                                                                                           /rpt_family="Aluy"
complement/innin
                                                                                                                                                                                                                                                                                                                                                                                                                /rpt_family="Aluy"
9409. .9695
                                                                                                                                                                                                                                                             /rpt
                                                                                                                                                                                                                                                                                                                                                     frame:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          frame: 2, quality: excellent(7290...7443)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reame: 1, quality: excellent, score: 93.000" complement(5378. .5511) /rpt_family="Almith"
                                                                                                                                                                                                                                                                                                                     rpt_family="Alusg/x"
                                                                                                                                                                                                                                                                                                                                           /note="predicted exon, program: grail2exons_human_1.3,
frame: 2, quality: excellent, score: 76.000"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /rpt_family="Alusg'complement(6285...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /rpt_family="L1MB7"
4230. .4530
                                                                                                                                                                                                                                                                                                                                                                                                rpt_family-"Alusg"
                                                                                                                                                                                                                                                                                                                                                                                                                                                           rpt_family="Aluy"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /rpt_family="BC200"
7472. .7588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  complement (5979.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /rpt_family="L1M4"
complement/gran
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rpt_family="AluJb"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rpt_family="FLAM_C"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /rpt_family="Alusx"
1663. .674"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /rpt_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 complement(5080.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         complement (4680
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /rpt_family="Alusx"
3866. .4229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rpt_fami.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rpt_family="Alusg1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'note="predicted exon,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       rpt_family-"MIR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /rpt_family-"FLAM_C"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rpt_family="L1pB3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'rpt_family-"Aluy"
                                                                                                                                                                                    _family="AluJo"
9. .21272
                                                                                                                                                                                                                                               _family="AluJ"
lement(1047)
                                                                          _family="AluSg"
                                                                                                       family="AluSx"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                family="Alusq"
ement(5307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             family-"AluSg/x
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         family-"L1MB7"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ly="AluSx"
.12772)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .6279)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .5079)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .4736)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         xon, program: grail2exons_human_1.3,
excellent, score: 97.000"
                                                                  Matches
                                                                                  Best Local
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Query Match
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      Similarity
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AA317001 EST188650 HCC cell line (metastasis to liver in mouse) II Homo sapiens cDNA 5' end similar to similar to ornithine decarboxylase antizyme (1.200); 1008
identity.—AA308557 EST179376 HCC cell line (matastasis to liver in mouse) II Homo sapiens cDNA 5' end similar to similar to ornithine decarboxylase antizyme (1.200); 1008
identity.—(14748.14942) AA307682 EST178550 Colon carcinoma (HCC) cell line Homo sapiens CDNA 5' end similar to similar to ornithine decarboxylase antizyme (1.195); 1008 identity.—(14758.14942) L49059 | HUMO10907F Homo sapiens retinal fovea EST HFD010907 sequence. (221.37); 1008 identity.—Other EST matches: T53070, T47553, T27358, T19460"
                                                                /note="DDS similarity to overlapping ESTs:
/note="DDS similarity to overlapping ESTs:
AA317001 EST188650 HCC cell line (metastasis to liver in mouse) II Homo sapiens cDNA 5' end similar to similar to ornithine decarboxylase antizyme (201. 346); 99% identity.-AA308557 EST179376 HCC cell line (matastasis to liver in mouse) II Homo sapiens cDNA 5' end similar to similar to ornithine decarboxylase antizyme (201. 346); 100% identity.-AA307682 EST178550 Colon carcinoma (HCC) cell line Homo sapiens cDNA 5' end similar to similar to
                                                                                                                                                                                                                                                                                                                                                                                                                             complement (16068. .16 /rpt_family="GC_rich"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="translation of first 68 residues of ODC antizyme; /note="translation of first 68 residues of ODC antizyme; Ornithine decarboxylase antizyme is likely expressed through translational frameshifting which results in the products of ORF1 and ORF2 being synthesized as a single polypeptide. In rat (D10706) the mRNA sequence at the junction of the two ORF5, UCC UGA U (253. .259, shown as ORF1 codons), is decoded as Ser-Asp through +1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /rpt_family="GC_rich"
complement(14613...14683)
/rpt_family="C"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SSSESSRVSLHCCSNPGPGPRWCS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /product="ODC antizyme ORF1"
/protein_id="AaCO2802.1"
/db_xref="G1:2894632"
/tzanslation="MYKSSLORILNSHCFAREKEGDKPSATIHASRTMPLLSLHSRGG
                                                                                                                                                                                                                                                                                                                                                                         /gene="OAZ"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (U09202) ornithine decarboxylase antizyme [Homo sapiens]."
join(14797. .14942,16583. .>16640)
/gene="OAZ"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /rpt_family="GC_rich"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 join(14797. .14942,16583. .16728,16980. .17152,17933. .18018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /codon_start=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /gene="OAZ"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18170. .18306)
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complement(14430...14
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13068. .13215
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complement(12872. .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note="ornithine decarboxylase antizyme; 98% identical to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /rpt_family="Alusx"
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5.1%; Score 19; DB 9; Length 37635; 100.0%; Pred. No. 16;
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35070 AAAAGGGGAAGTTCTGCCT 35088

317 AAAAGGGGAAGTTCTGCCT 335

Conservative

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Mismatches

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REFERENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Carron, T.F., Carter, M., Cavasos, S.R., Chacko, J., Chavez, D., Carron, T.F., Carter, M., Cavasos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Charko, D., Chavez, D., Cheveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., David, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., David, R., David, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., David, R., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwalte, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., Dearnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Ernandez, D., Harris, C., Harris, C., Harris, C., Harris, C., Harris, C., Harris, C., Harris, K., Hart, M., Havlak, P., Hale, S., Hamilton, K., Gorrell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K., Harris, C., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Jackson, L., E., Jackson, E., Homsi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E., Jackson, E., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C., Kratovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L., Liu, W., Loulseged, H., Lozdon, R.J., Liu, W., Loulseged, H., Lozdon, R.J., Lu, X., Lucier, A., Lucler, R., Luna, R., Ma, J., Mareshwari, M., Mapua, P., Martin, R., Martindale, A., Martinez, E., Manes, Y., Martindale, A., Martinez, E., Marer, S., Matchell, T., Mohabbat, K., Morgan, M., Morris, S., Moser, M., Norris, S., Matchell, T., Mohabbat, K., Morgan, M., Morris, S., Martine, E., Pu, L., Oulles, M., Ren, Y., Feters, L., Pickers, R., Pathus, E., Pu, L., Oulles, M., Ren, Y., Sodergen, E., Sonaike, T., Sparks, A., Stanley, H., Stose, H., Shosshtari, N., Stone, H., Shosshtari, N., Stone, H., Shosshtari, N., Stone, H., Shosshtari, N., Villalon, D., Vinson, R., Wang, Q., Wallalamson, A., Tamerisa, A., Tamerisa, K., Wang, Q., Wallalamson, A., Martinez, S., Warren, R., Washington, S., Wang, Q., Wan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Muzny, D.M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Bimage, K., Blankenburg, K., Bonnin, D., Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buhay, C., Burch, P., Burkett, C., Burrell, K.L., Byrd, N.C., Burhay, C., Burch, P., Burkett, C., Burrell, K.L., Byrd, N.C.,
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (19-JUL-2002) Human Genome Sequencing Center, Department
of Molecular and Human Genetics, Baylor College of Medicine, One
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Worley, K.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wu,C., Wu,Y., Wu,Y.F., Zhou,J., Weinstock,G. and Gibbs,R.
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                                                                 Center project Information Center project name: GZNK Center clone name: CH230-432K14
                                                                                                                                                                          Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
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*** SEQUENCING IN PROGRESS
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runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
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Homo sapiens BAC clone RP11-524B5
AC105316
AC105316.5 GI:205ססייר
                      Unpublished
                          Buatsi,D., Haakenson,W., Boye
The sequence of Homo sapiens
Unpublished (2001)
                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 148571)
Sulston,J.E. and Waterston,R.
Toward a complete human genome sequence Genome Res. 8 (11), 1097-1108 (1998)
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         Waterston, R.H.
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123175: contig of 10659 1
123275: gap of unknown 18
132799: contig of 9524 9
132899: gap of unknown 1
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                                                                                                                                                                                               Waterston, R.
Direct Submission
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Submitted (10-MAY-2002) Genome
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Center project name: H_NH0524B05
                                      Contact: sapiens@watson.wustl.edu
                                                        Center code: WUGSC
Web site: http://genome.wustl.edu/gsc
                                                                                           Center: Washington University Genome Sequencing Center
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NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions

This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest.

## MAPPING INFORMATION:

Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. Louis MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc

The RPCI-11 human BAC library was made from the blood of one male donor, as described by Oscegawa, K., Woon, P.Y., Zhao, B., Frengen, E. Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong Research Genetics, Inc. (http://www.reand coworkers at http://www.chori.org SOURCE INFORMATION: OR: pBACe3.6

# The clone sequenced to the left is RP11-620C21, 2000 bp overlap; the clone sequenced to the right is RP11-138A23, 2000 bp overlap Actual start of this clone is at base position 70882 of RP11-620C21; actual end is at base position 47389 of RP11-138A23 Location/Qualifiers

bp overlap

NEIGHBORING SEQUENCE INFORMATION:

FEATURES

source

repeat\_region /clone\_lib="RPCI-11" 138. .322 /db\_xref="taxon:9606" /chromosome="4" /map="4" /clone="RP11-524B5" /organism="Homo sapiens" .14857

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Local Similarity 100.0%; Pred. No.
                                                               Direct Submission
Submitted (11-NOV-2000) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St. Louis,
                                                                                                                                                                                                                  Homo sapiens.

Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 154693)

Sulston, J. E. and Waterston, R.

Toward a complete human genome sequence
Genome Res. 8 (11), 1097-1108 (1998)

99063792
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HOMO sapiens BAC clone RP11-215J17 from 4, complete sequence.
AC084750
AC084750.3 GI:17155079
Submitted (30-NOV-2001) Genome Sequencing Center, Washington
                   MO 63108, USA
4 (bases 1 to 154693)
Waterston, R.H.
Direct Submission
                                                                                                                                           2 (bases 1 to 154693)
Isak,A. and Kozlowicz,A.
The sequence of Homo sapiens BAC clone RP11-215J17
Unpublished (2001)
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Submitted (03-JAN-2002) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St. Louis,
MO 63108, USA
6 (bases 1 to 154693)
Waterston,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap
                                                                                                                                                                                                                                                                                               The clone sequenced to the left is RP11-362M19, 2000 bp overlap; the clone sequenced to the right is RP11-509I10. Actual start of this clone is at base position 151949 of RP11-362M19; actual end is at base position 154693 of RP11-215J17.
                                                                                                                                                                                                                                                                                                                                                                                                                             The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Frenger, E., Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at the Roswell Park Cancer Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MAPPING INFORMATION:
Mapping information for this clone was provided by Dr. John D.
Mapping information for this clone was provided by Dr. John D.
McPherson, Department of Genetics, Washington University, St. Louis
MO. For additional information about the map position of this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
                                                                                                                                                                                                                                           Data from AC091490 was used to finish this clone, AC084750 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             between neighboring data submissions.
                                                                                                                                                                                                                                                                                                                                                                           NEIGHBORING SEQUENCE INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                          (http://bacpac.med.buffalo.edu)
VECTOR: pBACe3.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOURCE INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Center project name: H_NH0215J17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: sapiens@watson.wustl.edu
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             /rpt_family="AT_rich" 6208. .6327
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      /rpt_
23610
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21358
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21164
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20805
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20262. .20282
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14936. .15017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /rpt_
14191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /rpt_family="L1"
11857. .11907
                                    /rpt_family="AT_rich" 32444. .32473
                                                                                                                                                                                                                                                                                                                                                                               /rpt_family="GA-rich" 25117. .26953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /rpt_family="Alu"
14487. .14548
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14115
                                                                                                                                                                                                                                                                                                       28399
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13902. .14148
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                                                                                                                                                                                                                              ....119="(TTTA)n"
.31192
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       .33269
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                                                                                                                                                                                                            -"MaLR"
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RESULT 32
AC106955
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VERSION
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Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AUTHORS
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                                                                                                                                                                                                                                              Muzny, D.M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Bimage, K., Blankenburg, K., Bonnin, D., Bouck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buhay, C., Burch, P., Burkett, C., Burrell, K.L., Byrd, N.C., Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, Z., Chow, D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Denn, A.L., Ding, Y., Dinh, H.H., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., Earnhart, C., Edgar, D., Flagg, N., Ford, J., Foster, P., Frantz, P., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Garcia, A., Garner, T., Garza, N., Gill, R., Harris, K., Hart, M., Havlak, P., Hale, S., Hamilton, K., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Harrandez, O., Hondroch, A., Havlak, P., Hale, S., Hamilton, R., Chille, R., Charles, M., Harlandez, J., Harlandez, J., Harlandez, D., Hades, A., Hernandez, J., Harlandez, J., Harlandez, D., Hades, A., Hernandez, J., Harlandez, J., Harland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ***, 7
                           Hernandez,O., Hodgson,A., Hogues,M., Holloway,C., Hollins,B.,
Homsi,F., Howard,S., Huber,J., Hulyk,S., Hume,J., Jackson,L.E.,
Jacobson,B., Jia,Y., Johnson,R., Jolivet,S., Joudah,S.,
Karlsson,E., Kelly,S., Khan,U., King,L., Korvah,J., Kovar,C.,
Krartovic,J., Kureshi,A., Landry,N., Leal,B., Lewis,L.C., Lewis,L.,
Li,J., Li,Z., Lichtarge,O., Lieu,C., Liu,J., Liu,W., Loulseged,H.,
Lozado,R.J., Lu,X., Lucier,A., Lucier,R., Luna,R., Ma,J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HTG; HTGS_PHASE1.
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Maheshwari,M., Mapua,P., Martin,R., Martindale,A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (bases 1 to 154772)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         55 154772 bp DNA linear HTG 13-JUL-2002 norvegicus clone CH230-119A22, *** SEQUENCING IN PROGRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /rpt_1
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43011
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44772. .44792
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44591
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43525
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; Pred. No.
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AUTHORS
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (13-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 12, 2002 this sequence version replaced g1:18701823.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (14-JAN-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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Direct Submission
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* This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequencing vector: Plasmid; Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap: version 0.990329 Consensus quality: 8851 bases at least Q40 Consensus quality: 93751 bases at least Q30 Consensus quality: 97432 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Center project name: GOOY Center clone name: CH230-119A22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Web site: http://www.hgsc.bcm.tmc.edu/
Contact: hgsc-help@bcm.tmc.edu
------Project Information
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* of the gaps between colors
* of the gaps between colors
* provided by the submittor.

* This sequence will be replaced
* by the finished sequence as soon as it is available and
* the accession number will be preserved.
* 1 2249: contig of 2249 bp in length
* 2250 2349: gap of unknown length
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On Aug 1, 2002 this sequence version replaced g1:22002191.
                                                                                                                                                                                                                  Center: Department Of Chemistry And Biochemistry
The University Of Oklahoma
Center code: UOKNOR
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                                                                                                                                                                                                                                                                                                    Direct Submission
Submitted (16-AUG-2002) Department Of Chemistry And Biochemistry, The University Of Oklahoma, 620 Parrington Oval, Room 208, Norman
                                                                                                                                                                                                                                                                                                                                                                                            Submitted (28-FEB-2002) Department Of Chemistry And Biochemistry, The University Of Oklahoma, 620 Parrington Oval, Room 208, Norman,
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87542:
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of 1939
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of 2194
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of 1772
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unknown
of 1680
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of 1714
of 1714
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of 2563
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12667 CTCCCCAGGGAAACTGAAC 12649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       source
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                                                                                                                                                                                                                  Submitted (26-JUN-2001) Sanger Centre, Hinxton, Cambridgeshire, CB10 18A, UK. E-mail enquiries: humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk
On Jun 27, 2001 this sequence version replaced gi:14272267.
During sequence assembly data is compared from overlapping clones.
Where differences are found these are annotated as variations together with a note of the overlapping clone name. Note that the variation annotation may not be found in the sequence submission corresponding to the overlapping clone, as we submit sequences with only a small overlap as described above.
This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one M13 subclone; and the assembly was confirmed by restriction digest. The following abbreviations are used to associate primary accession numbers given in the feature table with their source databases: Em:. EMBL; Sw:, SWISSPROT; Tr:, TREMBL; wp:, WORMPED; Information on the WORMPED than all property and the contract of the contract of the wormpen This sequence.
                                                                                                                                 http://www.sanger.ac.uk/Projects/C_elegans/wormpep This sequence was generated from part of bacterial clone contigs of human chromosome 9, constructed by the Sanger Centre Chromosome 9 Mappi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                           http://www.sanger.ac.uk/HGP/Chr9 RP11-145E5 is from the library RPCI-11.1 constructed by the group of Pieter de Jong. For further details see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
http://www.chori.org/bacpac/home.htm
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                                                                                                               Further information can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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/clone_lib="RPCI - 23 Female (C57BL/6J) Mouse BAC Library"
33372 c 32918 g 43471 t 500 others
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100.0%; Pred. No.
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om clone RP11-145E5 on chromosome
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                                                                                                                                          Chromosome 9 Mapping
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RESULT 35
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RRS Muzny, D. M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbrooks, S.L., Barch, M., Bryant, N.D., Bouck, J., Bouch, D., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buhay, C., Burch, P., Burkett, C., Burrell, K.L., Byrd, N.C., Carron, T. F., Carter, M., Cavazos, S.R., Chocko, J., Chavez, D., Chen, R., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Dathorne, S.R., David, R., Durbin, K.J., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Frantz, P., Falts, T., Ferraguto, D., Edwards, C.C., Elhaj, C., Escotto, M., Frantz, P., Falts, A., Garner, T., Garza, N., Gill, R., Gorrell, J., Ford, J., Foster, P., Frantz, P., Falts, A., Garner, T., Garza, N., Gill, R., Hernandez, J., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Hernandez, J., Hulyk, S., Hume, J., Jackson, L.E., Jacobson, B., Jak, Y., Johnson, R., Jollvett, S., Joudah, S., Kartovic, J., Kuresh, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L., Li, J., Liu, J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IMPORTANT: This sequence is not the entire insert of clone RR11-145E5 It may be shorter because we sequence overlapping sections only once, except for a 100 base overlap. The true left end of clone RR11-145E5 is at 1 in this sequence. The true left end of clone RR11-408N14 is at 157434 in this sequence. The true right end of clone RR11-149I2 is at 48466 in this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rattus norvegicus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence.
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AC116273.5 GI:21746202
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (18-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 14, 2002 this sequence version replaced gi:20467689.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (26-MAR-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 163041)
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Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                        NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a "working draft" sequence. It currently consists of 59 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence
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Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap: version 0.990329
Consensus quality: 116366 bases at least Q30
Consensus quality: 120870 bases at least Q30
Consensus quality: 124554 bases at least Q30
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------ Project Information
Center project name: GTXN
Center clone name: CH230-354C23
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Center code: BCM
Web site: http://www.hgsc.bcm.tmc.edu/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (25-APR-2000) Human Genomic Center, Institute of Genetics, Chinese Academy of Sciences, Datun Road, Beijing, Beijing 100101, P.R.China
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Center project name: 1% project
Center clone name: RP11-34405
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                                               Eukaryota; Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 177070)
Waterston, R.H.
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AC018875.7 GI:12963043
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The sequence of Homo sapiens clone
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Submitted (21-DEC-1999) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St. MO 63108, USA
On Feb 18, 2001 this sequence version replaced qi:11192196.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Insert size: 178000; agarose-fp
Insert size: 176607; sum-of-contigs
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Sequencing vector: plasmid; 14%
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Waterston, R.H.
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2 (bases 1
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                                                                  /note="assembly_name:Contig38" 35320. .36695
/note="assembly_name:Contig41"
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39649. .41067
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29937. .31395
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27246. .29836
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25939. .27145
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/note="assembly_name:Contig30"
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22712. .25838
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/organism="Homo sapiens"
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L5998. .17916
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L0332. .12457
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79153: gap of unknown length
109291: contig of 30138 bp in length
109391: gap of unknown length
177070: contig of 67679 bp in length
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Sequencing vector: MI3: 100%
Sequencing vector: MI3: 100%
Sequencing vector: plasmid: 0%
Sequencing vector: plasmid: 0%
Chemistry: Dye-primer ET: 100% of reads
Chemistry: Dye-terminator Big Dye: 0% of reads
Assembly program: Phrap; version 0.99039
Consensus quality: 154559 bases at least Q40
Consensus quality: 161004 bases at least Q30
Consensus quality: 166430 bases at least Q20
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Homo sapiens chromosome 11 clone RP11-567M21, WORKING DRAFT
SEQUENCE, 30 unordered pieces.
                                                                                                                                                                                                                   Center: Washington University Genome Sequencing Center Center code: WUGSC
                                                                                                                                                                                                                                                                                                         Direct Submission Submitted (15-JAN-2000) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St. MO 63108, USA
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Waterston, R.H.
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1 (bases 1 to 177426)
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47537. .51069
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44701. .47436
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109392. .177070
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79154. .109291
/note="assembly_name:Contig50
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51170. .56386
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Insert size: 189000; agarose-fp Insert size: 174526; sum-of-contigs Quality coverage: 2.77 in Q20 bases; agarose-fp Quality coverage: 3.06 in Q20 bases; sum-of-contigs
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* This record will be updated with the finished sequence.
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72619: gap of unknown length
79368: contig of 6749 bp in let
79468: gap of unknown length
88631: contig of 9163 bp in let
88731: gap of unknown length
98171: contig of 9440 bp in let
98271: gap of unknown length
108874: contig of 10603 bp in let
108974: gap of unknown length
110856: contig of 11882 bp in let
113832: contig of 12876 bp in let
            133932: gap of unknown length
151337: contig of 17405 bp in
151437: gap of unknown length
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FEATURES

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                                                                              /note="assembly_name:Contig40"
151438. 177426
/note="assembly_name:Contig41"
38792 c 37341 g 49554 t
                                                                                                                                                                                                                                                /note="assembly_name:Contig36"
98272. .108874
                                                                                                                                                                                                                                                                                      79469. .88631
/note="assembly_name:Contig35"
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                                                                                                                                                /note="assembly_name:Contig39"
133933. .151337
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                                                                                                                                                                              /note="assembly_name:Contig38"
120957. .133832
                                                                                                                                                                                                                          /note="assembly_name:Contig37"
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21851. .24673
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note-"assembly_name:Contigl9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note="assembly_name:Contig18"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="RP11-567M21"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note="assembly_name:Contig16"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="assembly_name:Contig13"
                5.1%; Score 19; DB 2; Length 177426; 100.0%; Pred. No. 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Alburoks, S. L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Binage, K., Blankenburg, K., Bonnin, D., Bouck, J., Bowie, S., Brévan, E., Brown, E., Brown, M., Bryant, N.P., Burky, C., Burch, P., Burkett, C., Burrell, K.L., Byrd, N.C., Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davis, C., Davy, Carroll, L., Dederich, D.A., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwalte, K.J., Delgado, D., Edwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Flagg, N., Ford, J., Foster, P., Frantz, P., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Flagg, N., Ford, J., Foster, P., Frantz, P., Foster, P., Frantz, P., Gorrell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K., Gorrell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K., Harris, C., Hartis, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Hernandez, O., Hodgson, A., Hogues, M., Holloway, C., Hollins, B., Honsi, F., Howard, S., Lin, Haylak, P., Hawes, A., Hernandez, J., Lin, F., Howard, S., Kratovic, J., Kureshi, A., Landry, N., Leal, B., Leets, J.C., Lewis, L.C., Liu, J., Liu, W., Loulseged, H., Lozado, R.J., Lu, X., Lucier, A., Lucler, R., Luna, R., Wartinez, E., Massey, E., Mawhiney, E., McLeod, M.P., Martindale, A., Martinez, E., Massey, E., Mawhiney, E., McLeod, M.P., Martindale, A., Martinez, E., Massey, E., Makerson, E., Wokenskwo, S., Oguh, M., Okwuonu, G., Oragunye, N., Oviedo, R., Parce, A., Payton, B., Peery, J., Perez, L., Pickens, R., Pinmus, E., Put., L., Oulles, M., Ren, Y., Perez, J., Put., E., Mosshari, K., Vasguez, L., Vera, V., Villalon, D., Vinson, R., Wang, Q., Wang, S., Walte, A., Tabor, P., Tamerisa, A., Tamerisa, S., Wang, Q., Wang, S., Walte, A., Tabor, P., Panerisa, A., Tanerisa, R., Tang, H., Stone, H., Stone, H., Wang, Q., Walte, M., S., Sherrer, S., Shen, H., Shooshari, N., She
Submitted (17-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One
                                                                                                                                                                                                            Submitted (21-FEB-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                  Worley, K.C.
                                                                                                                                                                                                                                                                                                                               Worley, K.C.
Direct Submission
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AC112477
AC112477.3 GI:21745567
HTG; HTGS_PHASE1.
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***, 63 unordered pieces.
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COMMENT

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Baylor Plaza, Houston, TX 77030, USA on Jul 14, 2002 this sequence version replaced gi:20303368.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   * NOTE: Estimated insert size may differ from sequence length
(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).
*NOTE: This is a 'working draft' sequence. It currently
* consists of 63 contigs. The true order of the pieces
* is not known and their order in this sequence record is
* arbitrary. Gaps between the contigs are represented as
* runs of N, but the exact sizes of the gaps are unknown.
* This record will be updated with the finished sequence
* as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     as soon as it is available be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Center project name: GSSY
Center clone name: GH30-191H11
Center clone name: CH30-191H11
Center clone name: CH30-191H11
Center clone name: CH30-191H11
Center clone vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329
Consensus quality: 130102 bases at least Q40
Consensus quality: 143613 bases at least Q30
Consensus quality: 148883 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Web site: http://www.hgsc.bcm.tmc.edu/
Contact: hgsc-help@bcm.tmc.edu
------Project Information
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Center: Baylor College of Medicine
Center code: BCM
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       RESULT 40
                                                Db 42662 GAAGTGATGGAGCAAAGTG 42680
                                                                                                       Query Match
Best Local Similarity
Matches 19; Conser
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                            Birren, B., Linton, L., Nusbaum, C., Lander, E., Ali, A., Allen, N., Anderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B., Anderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B., Brown, A., Chang, J., Chazaro, B., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cook, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Faro, S., Ferreira, P., Fitzhugh, W., Gage, D., Galagan, J., Gardyna, S., Glode, S., Gord, S., Goyette, M., Graham, L., Grand-Plerre, N., Hagos, B., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, A., Karatas, A., Kells, C., LaRocque, K., Lamazares, R., Landers, T., Lehoczky, J., Levine, N., Liu, G., MacLean, C., Landers, T., Lehoczky, J., Levine, N., Liu, G., MacLean, C., Landers, T., Lehoczky, J., Levine, N., Landers, C., MacLean, C., Landers, T., Lehoczky, J., Levine, N., Landers, C., MacLean, C., Landers, T., Lehoczky, J., Levine, N., Landers, C., MacLean, C., Landers, T., Lehoczky, J., Levine, N., Landers, C., MacLean, C., Landers, T., Lehoczky, J., Levine, N., Landers, C., MacLean, C., Landers, C., Land
Macdonald, P., Major, J., Marquis, N., Matthews, C., McCarthy, M.,
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On Aug 9, 2002 this sequence version replaced gi:20334333.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fujlyama, A., Yada, T., Totoki, Y., Watanabe, H. and Sakaki, Y. Homo sapiens genomic DNA Published Only in Database (2000) 2 (Dases 1 to 188172) 4 (Dases 1 to 188172) 4 (Dases 1 to 184172) 5 (Dases 1 to 184172) 7 (Dases 1 to 184172) 7 (Dases 1 to 184172) 8 (Dases 
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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COMMENT

This sequence will be replaced by the finished sequence as soon as it is available and

the accession number will be preserved

TITLE

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REFERENCE
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* consists of 9 contigs. Gaps between the contigs are represented as runs of N. The order of the pieces is believed to be correct as given, however the sizes of the gaps between them are based on estimates that have provided by the submittor.
                                                                                                                                                                                                                                 * NOTE: This is a 'working draft' sequence. It currently
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Smit, A.F.A. & Green, P. (1996-1997)
http://ftp.genome.washington.edu/RM/RepeatMasker.html
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                                                                                                                                                                                                                                                                                                                      Summary Statistics

Sequencing vector: Plasmid; n/a; 100% of reads
Chemistry: Dye terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.960731
Consensus quality: 191646 bases at least Q40
Consensus quality: 193107 bases at least Q30
Consensus quality: 193517 bases at least Q20
Insert size: 194000; agarose-fp
Insert size: 193969; sum-of-contiss
Quality coverage: 6.0 in Q20 bases; agarose-fp
Quality coverage: 6.0 in Q20 bases; sum-of-contigs
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Center clone name: 358_A_18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: sequence_submissions@genome.wi.mit.edu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zainoun, J., Zembek, L., Zimmer, A. and Zody, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   320 Charles Street, Cambridge, MA 02141, USA
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FEATURES

source

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TITLE JOURNAL

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68166 ACTCAGTTTGCATCAGCTG 68148
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                                                                                                                                                                                                                                                                     AC122259 209418 bp DNA linear HTG 23-1
Mus musculus chromosome UNK clone RP23-183E7, WORKING DRAFT
SEQUENCE, 10 unordered pieces.
AC122259
AC122259.1 GI:21105101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 1980: contig of 1980 bp in length
2081 3501: gap of 100 bp
3502 3601: gap of 100 bp
6848: contig of 3247 bp in length
6849 6948: gap of 100 bp
11585 11684: gap of 100 bp
11685 30031: contig of 4636 bp in length
30032 30131: gap of 100 bp
30132 88538: contig of 18347 bp in length
18853 88638: gap of 100 bp
188539 88638: gap of 100 bp
118825: contig of 58407 bp in length
118826 118925: gap of 100 bp
118926 162363: contig of 33187 bp in length
162364 194769: contig of 33306 bp in length
162464 194769: contig of 33306 bp in length
162463: gap of 100 bp
162464 194769: contig of 33306 bp in length
162464 194769: contig of 33306 bp in length
                                                                                                1 (bases 1 to 209418) McPherson, J.D. and Waterston, R.H.
                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                            Mus musculus
                                            Unpublished
                                                                    The sequence of Mus musculus
                                                                                                                                                                                                                                                HTG; HTGS_PHASE1; HTGS_DRAFT
McPherson, J.D. and Waterston, R.H.
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                    (bases 1 to 209418)
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38216 c 39235 g
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118926. .162363
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequencing vector: M13; 0%
Sequencing vector: M13; 0%
Sequencing vector: plasmid; 100%
Chemistry: Dye-primer ET; 0% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.990319
Consensus quality: 205472 bases at least Q40
Consensus quality: 205472 bases at least Q30
Consensus quality: 2054860 bases at least Q20
Insert size: 212000; agarose-fp
Insert size: 208518; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Direct Submission Submitted (23-MAY-2002) Genome Sequencing Center, 4444 Forest Park Parkway, St. Louis, MO 63108, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Center project name: M_BA0183E07
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Contact: submissions@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         * NOTE: This is a 'working draft' sequence. It currently consists of 10 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Quality coverage: 7.49 in Q20 bases; agarose-fp Quality coverage: 6.37 in Q20 bases; sum-of-contigs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     * arbitrary. Gaps between the contigs are represented as
* runs of N, but the exact sizes of the gaps are unknown.
* This record will be updated with the finished sequence
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88783
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1923447
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193493
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6583
19748
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19848
51249
                                                                                                                                                                            /note="assembly_name:Contigl2"
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125736. .192346
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/chromosome="UNK"
                                                                                                                                                                                                                                                                                                                                                                                                                                                     1. .209418
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  location/Qualifiers
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                                                                                                                                                                                                                                                                                                     note="assembly_name:Contig10"
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19747:
19847:
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2: gap of unknown length
7: contig of 13165 bp in length
7: gap of unknown length
8: contig of 31401 bp in length
8: gap of unknown length
2: contig of 37334 bp in length
2: contig of 36833 bp in length
5: contig of 66611 bp in length
6: contig of 66611 bp in length
6: contig of 1046 bp in length
6: gap of unknown length
6: gap of unknown length
8: gap of unknown length
9: contig of 5327 bp in length
9: gap of unknown length
9: contig of 5327 bp in length
9: gap of unknown length
9: gap of unknown length
9: contig of 5327 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gap of unknown contig of 5803
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SOURCE ORGANISM

KEYWORDS VERSION ACCESSION DEFINITION

REFERENCE

REFERENCE

AUTHORS

JOURNAL TITLE AUTHORS RESULT 42 AC122259

В S BASE COUNT ORIGIN

misc\_feature misc\_feature misc\_feature misc\_feature misc\_feature misc\_feature misc\_feature

misc\_feature

Query Match Best Local

Matches

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Db 134911 GCCAAGGCCCAAGAAAAGG 134893
                                                                                                                                                                                                                     BASE COUNT
ORIGIN
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Best Local :
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                                                                                                                                                             Local Similarity
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eature 203616. 209418
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58209 a 45087 c 44929 g 60292 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (23-OCT-2001) DOE Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA
On Oct 23, 2001 this sequence version replaced gi:7711412.
Draft Sequence Produced by DOE Joint Genome Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (15-SEP-1999) Production Sequencing Facility, DOE Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA (bases 1 to 238472)
DOE Joint Genome Institute and Stanford Human Genome Center.
                                                                                                                                                                                                                                                                                                                                                                     Quality: Phrap Quality >=40 99.7% of Sequence; Estimated Total Number of Errors is 0.5.
                                                                                                                                                                                                                                                                                                                                                                                                        www-shgc.stanford.edu
                                                                                                                                                                                                                                                                                                                                                                                                                          Finishing Completed at Stanford Human Genome Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Direct Submission
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DOE Joint Genome Institute.
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1 (bases 1 to 238472)
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/chromosome="5"
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                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
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192447. .193492
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Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
                                   Center code: BCM
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Mizny, D.M., Adins, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbarla, J., Benton, J., Binage, K., Blankenburg, K., Bonnin, D., Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buhay, C., Burch, P., Burkett, C., Burcell, K.L., Byrd, N.C., Cleveland, C.D., Cox, C., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M. D., Dathorne, S.R., David, R., Delancy, K.R., Davis, C., Davy-Carroll, L., Dederich, D.A., Delancy, K.R., Davis, C., Davy-Carroll, L., Dederich, D.A., Delancy, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinhi, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Flagy, M., Ford, J., Foster, P., Frantz, P., Falls, T., Forter, H., Dugan-Rocha, S., Durbin, K.J., Davis, C., Hamilton, K.J., Garcia, A., Garner, T., Garza, N., Gill, R., Gorrell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K., Harris, C., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Homards, C., Hodgson, A., Hogues, M., Holloway, C., Hollins, B., Homsi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Kalay, L., Kan, U., King, L., Kroth, J., Kovar, C., Karlsson, E., Kelly, S., Khan, U., King, L., Kroth, J., Kovar, C., Karlsson, E., Kally, N., Lucier, R., Luna, R., Ma, J., Lu, X., Lucier, A., Lucier, R., Luna, R., Martinez, E., Maheshwari, M., Mapua, P., Martin, R., Mardindle, A., Martinez, E., Maheshwari, M., Mapua, P., Martin, R., Mardindle, A., Martinez, E., Maheshwari, M., Mapua, P., Martin, R., Mardindle, A., Martinez, E., Petty, J., Petez, L., Peters, M., Mohabbat, K., Moygan, M., Morris, S., Moser, M., Roi, S., Norelson, S., Walland, R., Stanley, H., Stone, H., Shoth, R., Parks, A., Tamerisa, A., Tamerisa, K., Tang, H., Maris, G., Wall, H., Marting, G., Wall
                                                                                                                       Submitted (11-JUL-2002) Human Genome Sequencing of Molecular and Human Genetics, Baylor College Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                               Worley, K.C.
Direct Submission
Submitted (17-SEP-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
3 (bases 1 to 239904)
                                                                                                                                                                                                                                                                                                Worley, K.C
                                                                               On Jul 10, 2002 this sequence version replaced gi:18846093
                                                                                                                                                                                                                                                            Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA linear HTG 11-JUL-2002 Rattus norvegicus clone CH230-162P21, *** SEQUENCING IN PROGRESS ***, 44 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (bases 1 to 239904)
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Center: Baylor College of Medicine
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                                       Genome Center
                                                                                                                                                                   of Medicine, One
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NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 44 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Center project name: GEXO
Center clone name: CH230-162P21
Center clone name: CH230-162P21
Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of read Assembly program: Phrap; version 0.990329
Consensus quality: 19179 bases at least Q40
Consensus quality: 196736 bases at least Q30
Consensus quality: 201345 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    as soon as it is available and the accession number will be preserved.
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1 (bases 1 to 247865)

Muzny,D.M., Adams,C., Adio-Oduola,B., Ali-osman,F.R., Allen,C., Alstrooks,S.L., Amaratunge,H.C., Are,J.R., Ayele,M., Banks,T., Barbaria,J., Benton,J., Bimage,K., Blankenburg,K., Bonnin,D., Bouck,J., Bowie,S., Brieva,M., Brown,E., Brown,M., Bryant,N.P., Buthay,C., Burch,P., Burkett,C., Burerll,K.L., Byrd,N.C., Carron,T.F., Carter,M., Cavazos,S.R., Chacko,J., Chavez,D., Chen,G., Chen,R., Chen,Z., Chowdhry,I., Christopoulos,C.,
                                                                                                                                                                                                                                      247865 bp
Rattus norvegicus clone CH230-2G16,
40 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
19; Conser
                                                                                                                                                     Eukaryota; Metazoa;
                                                                                                                                                                     Rattus norvegicus
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/db_xref="taxon:10116"
/clone="CH230-162P21"
/49743 c 49983 g 64401 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                       Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          154817: gap of unknown length
164405: conclig of 9588 bp in length
164505: gap of unknown length
170277: conclig of 5772 bp in length
170377: gap of unknown length
179323: contig of 8946 bp in length
179423: gap of unknown length
179423: gap of unknown length
193498: contig of 14075 bp in length
193598: gap of unknown length
193598: gap of unknown length
208152: contig of 14554 bp in length
208252: gap of unknown length
223131: contig of 14879 bp in length
233231: gap of unknown length
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87909: contig

87909: gap of

9461: contig

94561: gap of

101192: contig

101292: gap of

107536: gap of

107536: gap of

11283: contig

11283: contig

11283: gap of

120978: gap of

120978: gap of

133119: contig

133119: contig

133219: gap of

143748: gap of

143748: gap of
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                                                                                                                                                                                                                                                                                                                                                                                                                             5.1%; Score 19;
100.0%; Pred. No.
                                                                                                                                 Chordata; Craniata; Vertebrata; Euteleostomi; Rodentia; Sciurognathi; Muridae; Murinae;
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SEQUENCING IN PROGRESS ***,
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Cleveland,C.D., Cox,C., Coyle,M.D., Dathorne,S.R., David,R.,
Davila,M.L., Davis,C., Davy-Carroll,L., Dederich,D.A.,
Delandey,K.R., Delgado,O., Denn,A.L., Ding,Y., Dinh,H.H.,
Douthwaite,K.J., Draper,H., Dugan-Rocha,S., Durbin,K.J.,
Earnhart,C., Edgar,D., Edwards,C.C., Elhaj,C., Escotto,M.,
Falls,T., Ferraguto,D., Flagg,N., Ford,J., Foster,P., Frantz,P.,
Gablsi,A., Gao,J., Garcia,A., Garner,T., Garza,N., Gill,R.,
Gorrell,J.H., Guevara,W., Gunaratne,P., Hale,S., Hamilton,K.,
Hernandez,O., Hodgson,A., Hogues,M., Holloway,C., Hollins,B.,
Homsi,F., Howard,S., Huber,J., Hulyk,S., Hume,J., Jackson,L.E.,
Jacobson,B., Jia,Y., Johnson,R., Jolivet,S., Joudah,S.,
Karlsson,E., Kelly,S., Khan,U., King,L., Korvah,J., Kovar,C.,
Karlsson,E., Kelly,S., Khan,U., King,L., Korvah,J., Kovar,C.,
Kratovic,J., Kureshi,A., Landry,N., Leal,B., Lewis,L.C., Lewis,L.,
Li,J., Li,Z., Licher,A., Lucier,R., Liu,J., Liu,J., Liu,J.,
Liu,X., Lucier,A., Lucier,R., Luna,R., Ma,J.,
Lay, Lu,X., Lucier,A., Face,C., Metador,M., Mei,G., Metzker,M.,
Maneshwari,M., Mapua,P., Martin,R., Martindale,A., Martinez,E.,
Manesey,E., Mavhiney,E., McLeod,M.P., Meador,M., Mei,G., Metzker,M.,
Miner,G., Miner,Z., Mitchell,T., Mohabbat,K., Morgan,M., Morris,S.,
Moser,M., Neal,D., Newtson,J., Newtson,N., Nguyen,A., Nguyen,N.,
Nguyen,N., Nickerson,E., Paylon,B., Peery,J.,
Petezs,L., Pickens,R., Primus,E., Pu,L.L., Quiles,M., Ren,Y.,
Rodergren,E., Sonaike,T., Sparks,A., Tamerisa,A., Tamerisa,N., Thomas,S.,
Tansey,J., Taylor,C., Taylor,T., Telfrod,B., Thomas,N., Thomas,S.,
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On Jul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (24-AUG-2002) Human Genome Sequencing Center, Depart of Molecular and Human Genetics, Baylor College of Medicine, Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (11-OCT-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 247865)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Worley, K.C.
Direct Submission
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                      (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 40 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence.
                                                                                                                                                                                                                                                                                                                                       NOTE: Estimated insert size may differ from sequence length
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                                                                                                                                                                                                                                                                                                                                                                                                              Center project name: TUUI
Center clone name: CH330-2016
Center clone name: CH330-2016
Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 211853 bases at least Q40
Consensus quality: 215254 bases at least Q30
Consensus quality: 217501 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Center code: BCM Web site: http://www.hgsc.bcm.tmc.edu/
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soon as it is available and the accession number will
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REFERENCE AUTHORS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /SIDS2/gcgdata/geneseq/geneseqn-embl/NA2000.DAT:*
                                                                                                                                                                                                                          Length
  369
1524
1619
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ABK33543
ABT03277
ABT03281
ABL40345
ABL40349
AAF93845
                                                                                                                                    ABT03274
ABL48956
                                                                                                                                                                                AAF95007
                                                                                                                                                                                                                                                                                              SUMMARIES
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                                                                                                                                                                                                                          Description
                                                                                                                                Human ovarian canc
Human ovarian carc
Ovarian carcinoma
                                                                   cDNA encoding huma
Human ovarian carc
Human ovarian carc
    Ovarian carcinoma
Ovarian carcinoma
Human cDNA encodin
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prosopnia meranog	ABL15972	23	8268	4.9	18	45
	14	23	6399	4.9	18	44
_	AAH54174	22	4106	4.9	18	43
. epidermidis	494	22	3633	4.9	18	42
uman polypucie	AAI59883	22	3556	4.9	18	41
s ge	AAH54960	22	3281	4.9	18	40
	AAS91414	23	2253	4.9	18	39
ine G713 encod	AAA55968	21	1791	-	18	38 8
ORFX	AAC74983	21	1633		18	37
Staphylococcus epi	ABN92620	24	1596		18	ω 6
ep	ABN91962	24	1587		18	S S
	AAS83729	23	1542	4.9	18	34
	AAH52496	22	1152		18	ယ
	AAK66340	22	484		18	32
immune/	AAK66339	22	484		18	31
Corn tassel-derive	ABL75647	24	286		18	30
	ABL76335	24	256		18	29
	ABL75913	24	237	4.9	18	28
Human cDNA encodin	ABK62081	24	2222	5.1	19	27
Human cDNA encodin	ABK62080	24	2080	5.1	19	26
Human reproductive	AAL07456	22	1226	5	19	25
Human ovarian canc	ABL81273	24	430	4	55	24
	ABL40351	24	480	_	117	23
Human ovarian carc	ABT03283	24	480	31.7	117	22
ant	ABQ54231	24	1608	8	142	21
10	AAF94044	22	591	9.	221	20
Ovarian carcinoma	ABL40350	24	1010	ω	271	19
Human ovarian carc	ABT03282	24	1010	ω ·	271	18
Ovarian carcinoma	ABL48768	24	396		281	17
Human ovarian carc	ABT03085	24	396	76.2	281	16
	AAF94818	22	396	5	281	15
Ω	AAF64188	22	1956	5	283	14
	AAF22400	21	1953	76.7	283	13
	AAD18690	22	2528		314	12
carcinon	ABL40352	24	1897	85.4	315	11
<	ABT03284	24	1897	85.4	315	10

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## ALIGNMENTS

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RESULT 1
AAF95007
ID AAF9
  10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                               Human ovarian cancer associated coding sequence SEQ ID NO: 199
                                                                                                                                                                                                                                                     AAF95007;
                                                                                                                                                                                                                                                                       AAF95007 standard; DNA; 369 BP
Isolated polypeptides associated with ovarian carcinomas, and the
                                                                                                                       08-SEP-2000; 2000WO-US24827
                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                               Human, ovarian
                                                                                                                                                                                                                                   23-MAY-2001 (first entry)
                   WPI; 2001-211395/21.
                                       Χu J,
                                                                                                                                         15-MAR-2001
                                                                                                                                                           WO200118046-A2
                                                        (CORI-) CORIXA CORP.
                                       Stolk JA;
                                                                          99US-0394374.
2000US-0561778.
2000US-0640173.
2000US-0656668.
                                                                                                                                                                                                cancer; vaccine; gene therapy; carcinoma; ds.
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RESULT 2
ABT03274
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Best Local
                              14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
         (CORI-) CORIXA CORP
                                                                                13-NOV-2001; 2001WO-US45395
                                                                                                           23-MAY-2002
                                                                                                                                  WO200239885-A2
                                                                                                                                                                                       Human; ovarian cancer; ovarian carcinoma; gene therapy;
                                                                                                                                                                                                               Human ovarian carcinoma associated coding sequence SEQ ID NO: 199.
                                                                                                                                                                                  cytostatic; gene;
                                                                                                                                                                                                                                            05-SEP-2002 (first entry)
                                                                                                                                                                                                                                                                                             ABT03274 standard; cDNA; 369
                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                         ABT03274;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention provides a number of coding sequences and proteins, the over-expression of which is associated with ovarian carcinoma/Cancer. These can be used in the diagnosis, treatment and prevention of ovarian cancer, optionally by gene therapy or in the form of a vaccine. The present sequence is an example of one of these sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 369 BP; 82 A; 109 C; 90 G; 86 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 18; Page 189; 189pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleic acids that encode them, useful for the prevention diagnosis and treatment of ovarian cancers - % \left( 1\right) =\left\{ 1\right\} 
                                                                                                                                                                                                                                                                                                                                                          361 CACCATCCT 369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           369;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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Best Local :
10-SEP-1999;
                       03-APR-2001; 2001US-0825294.
                                                                                                                                  Ovarian cancer; cancer therapy; vaccine; gene therapy; tumour; cancer;
                                                  10-JAN-2002
                                                                            US2002004491-A1
                                                                                                    Homo sapiens
                                                                                                                                                                Ovarian carcinoma sequence isolate 57887.
                                                                                                                                                                                          18-JUN-2002 (first entry)
                                                                                                                                                                                                                      ABL48956;
                                                                                                                                                                                                                                           ABL48956 standard; cDNA; 369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 369 BP; 82 A; 109 C; 90 G; 86 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer
                                                                                                                                                                                                                                                                                                             361
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                                                                                                                                                                                                                                                                                                                                    361 CACCATCCT 369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA 300
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                                                                                                                                                                                                                                                                                                            CACCATCCT 369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCGAGTTCATTGTGAATTGCAC
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99US-0394374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 369; DB 24; Length 369; 100.0%; Pred. No. 5.5e-187;
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                                                                                                                                                                                                                                                                                                                                                                                  Matches 369;
                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ovarian carcinoma proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ovarian tumour polypeptide and prevention and/or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ovarian tumour
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                         CACCATCCT 369
                                                CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCGGCAC
                                                                                                   GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA
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                                                                                                                      CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
                                                                                                                                                                                                                                                                 GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
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 CACCATCCT
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) ALGATE P A.
) FLING S P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Stolk JA, Algate PA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                     369 BP; 82 A; 109 C;
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                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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2000US-0640173.
2000US-0656668.
2000US-0713550.
369
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                                                                                                                                                                                                                                                                                                                                                                                            Score 369; DB 24;
Pred. No. 5.5e-187;
                                                                                                                                                                                                                                                                                                                                                                                                                                    90 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polynucleotide useful in diagnosis, cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                     86 T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     2 other;
                                                                                                                                                                                                                                                                                                                                                                                                         Length
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ABK33543
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01-DEC-2000;
20-DEC-2000;
20-DEC-2000;
20-DEC-2000;
28-FEB-2001;
10-MAY-2001;
25-MAY-2001;
                                                                                                                                                                                                                                                                            25-JUL-2000;
26-JUL-2000;
28-JUL-2000;
23-AUG-2000;
23-AUG-2000;
15-SEP-2000;
10-NGV-2000;
                                                                                                                                                                                                                                                                                                                               25-JUL-2000;
25-JUL-2000;
25-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; secreted process, forcest tumour; rectal tumour, rectal tumour, persicyte cell proliferation; chondrocyte cell proliferation; tumour necrosis factor-alpha; gene; ss.
                                                                                                                                                                                                           (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA encoding human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK33543 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                            25-JUL-2000;
25-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                     25-JUL-2000;
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2000WO-US23522
2000WO-US2332B
2000US-000000P
2000WO-US30873
2000WO-US3267B
2000WO-US3267B
2000WO-US3267B
2000WO-US3267B
2000WO-US3267B
2000WO-US3267B
2000WO-US34956
2001WO-US34956
2001WO-US34956
2001WO-US34280
2001WO-US34280
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2000US-220666P.
2000US-220893P.
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2000US-220638P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         colon cancer;
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Baker KP, Grimaldi 2002-172001/22. 'n, Desnoyers JC, Gurney AL, Gerritsen Smith V, Stephan Goddard A, Godows han JF, Watanabe Godowski PJ; Wood WI;

P-PSDB; AAU83599.

One hundred and twenty two nucleic acids encoding PRO polypeptides, useful for treating a PRO related disorder and for diagnosing tumours such as lung cancer, colon cancer, breast tumour, prostate tumour, rectal tumour or liver tumour.

Claim 2; Figure 15; 359pp; English.

The invention relates to one hundred and twenty two nucleic acids encoding PRO polypeptides. The sequences of the 122 PRO polypucleotides encode human secreted proteins. The PRO nucleic acids, polypeptides, agonists and antagonists are useful for treating a PRO related disorder. The PRO polypeptides are useful for diagnosing tumours, especially lung cancer, colon cancer, breast tumour, prostate tumour, rectal tumour or liver tumour. The PRO polypeptides are useful for stimulating the proliferation of, or gene expression, in pericyte cells, for stimulating the proliferation or differentiation of chondrocyte cells, for stimulating the release of tumour necrosis factor-alpha from human blood, for stimulating or inhibiting the proliferation of normal human dermal fibroblast cells. The PRO polypeptide may also be used as molecular weight markers and for tissue typing. The PRO nucleic acids have applications in molecular holder. in molecular biology, including use as hybridisation probes

RESULT 4

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RESULT 5
ABT03277
ID ABT03277
XX ABT03277
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03-APR-2001;
02-OCT-2001;
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                                                                                                        Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                                                                                                                                                                                              Xu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; ovarian cancer; cytostatic; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-SEP-2002 (first entry)
  The present invention provides human ovarian
                                               Claim
                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                          13-NOV-2001; 2001WO-US45395
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABT03277 standard; cDNA;
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                                                                                                                                                                                                                                                                                       (CORI-) CORIXA CORP
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les 315; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in chromosome and gene mapping. \mathtt{ABK33536}\text{-}\mathtt{ABK33657} represent human protein coding sequences of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCATCAGCTGCTACAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGAAGTT
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                                                                                                                                                                                                     2002-500186/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ovarian
                                             2;
                                                                                                                                                                                                                                            Stolk JA, Algate
                                               Page 189-190;
                                                                                                                                                                                                                                                                                                                                  2000US-0713550.
2001US-0825294.
2001US-0970966.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy;
         cancer
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      associated
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                                                                                                                   ovarian
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RESULT 6
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Best Local Similarity
Matches 315; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and coding sequences. The sequences can be used in the diagnosis treatment of ovarian cancers. The present sequence is a coding se
                                                                                                                                                                                                                                                      Human; ovarian cancer;
cytostatic; gene; ss.
                                                                                                                                                                                                                                                                                 Human ovarian carcinoma associated coding sequence SEQ ID NO: 211.
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                                   Novel ovarian cancer polypeptide and polynucleotide, detecting the presence of ovarian cancer in a patient pharmaceutical compositions, e.g. vaccines, for treat
                                                                            WPI;
                                                                                                                                    14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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                                                                                                                                                                                                                  WO200239885-A2
                                                                                              Xu J,
                                                                                                                                                                          13-NOV-2001; 2001WO-US45395
                                                                                                                                                                                               23-MAY-2002
                                                                                                                 (CORI-) CORIXA CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                                                                                                                                                                                                                                                                                                                    CTGCCTCGGCCCTCA 344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT
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                                                                                                                                                                                                                                    sapiens
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                                                                             2002-500186/53.
                                                                                              Stolk JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                               Algate
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                                                                                                                                                                                                                                                                 ovarian carcinoma; gene therapy; immunotherapy;
                                                                                                                                                                                                                                                                                                                                              1619
                                                                                                PA,
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Pred No. 4.16
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                                                                                                                                                                                                                                                                                                                                             ВP
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4.1e-158;
                                                                                                 Molesh
                                      a patient, and in for treating ovarian
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Claim

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Page 195; 197pp; English

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RESULT 7
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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                      (XUJJ/)
(STOL/)
(ALGA/)
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                                                                                                                                                                                                                    US2002004491-A1
                                                                                                                                                                                                                                                                       Ovarian
                                                                                                                                                                                                                                                                                         Ovarian carcinoma sequence isolate 57887 extended cDNA
                                                                                                                                                                                                                                                                                                               28-JUN-2002 (first entry)
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                                                                                                                                                                            03-APR-2001;
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                    WPI; 2002-171027/22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present invention provides human ovarian cancer associated proteins coding sequences. The sequences can be used in the diagnosis and atment of ovarian cancers. The present sequence is a coding sequence
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                                                            STOLK J A.

ALGATE P A.

FLING S P.
tumour polypeptide
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                                                                                                              2000US-0561778.
2000US-0640173.
2000US-0656668.
2000US-0713550.
                                                                                                                                                                             2001US-0825294
                                                                                                                                                        99US-0394374
                                                                                                                                                                                                                                                                       cancer therapy; vaccine; gene therapy; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   85.4%; Score 315; DB 24; 100.0%; Pred. No. 4.1e-158
  and
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  polynucleotide useful in
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the expanding T-cells specific for a tumour protein, and for inhibiting the
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                                                                                                                                                                                        Ovarian carcinoma O591S nucleotide sequence.
                                                                                                   Homo sapiens
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                                                                                                                                                         therapy; tumour; cancer;
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10-JAN-2002.

03-APR-2001; 2001US-0825294

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Comparish the invention are useful for their completent of the invention are useful for their continues of the invention are useful for their continues of the entire desired gene or gene fragments, and for designing and continues in tumour cells. Polypeptides and polynucleotides of the current comparish of a polypeptide in appropriate host cells. The current conspired the companies to direct convention are also useful in recombinant DNA molecules to direct convention are also useful in appropriate host cells. The current converged to the companies of the current converged to the cur
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AAF93845 standard; cDNA; 1890 BP
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01-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-171027/22
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2000US-0656668.
2000US-0713550.
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Pred. No. 4.1e-158;
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                                                                                                                                                                                                                                            Matches 315;
                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                            (e.g. polymerase chain reactions (PCR)) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and activity. The antibodies and antagonists may also be used as therapeutic agents to down regulate expression and activity. The antibodies and antagonists may also be used as diagnostic agents for detecting the presence of the polypeptides in samples (e.g. by enzyme linked immunosorbant assay (ELISA). Examples of diseases which may be treated include rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diagnosis of diseases associated with inappropriate secretory protein/membrane protein expression. The nucleic acids and complementary sequences may also be used as DNA probes in diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to nucleic acid sequences AAF93744 - AAF93916 which encode human secretory or membrane proteins represented by AAB88317 - AAB88419. Included in the invention are primers AAF93917 - AAF94295 and AAF62232 - AAF62235 which are used to isolate the cDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and
                                                                                                                                                                                                                                        Sequence 1890 BP; 419 A; 528 C; 533 G; 410 T; 0 other;
                                                                                                                                                                                                                                                                                  arthritis and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding secretory proteins/membrane proteins, useful gene therapy or as candidate target molecules in drug development -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P-PSDB; AAB88418
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      383
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                                                                              323 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 382
                                                                                                                    30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                                                                                                  Local Similarity
                    GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAAAGGTTCAAGAACATGTGTCAGAAAG
GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID 203; 609pp + CD ROM; English
                                                                                                                                                               Conservative
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2000JP-0118775.
2000JP-0183766.
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                                                                                                                                                                            85.4%; Score 315; DB 22; 100.0%; Pred. No. 4.2e-158;
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                                                                                                                                                                                               Length 1890;
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150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT

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RESULT 1
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03-APR-2001;
02-OCT-2001;
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                                                                                                                                                                                                                  Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                                                                                                                                                                                                                                                                                                             13-NOV-2001; 2001WO-US45395
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                                                                                                                                                                                                                                                                                                                                                                                                     WO200239885-A2
                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                            cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human ovarian carcinoma associated coding sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-SEP-2002 (first entry)
                                                                                                                                                The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
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                                                                                                                    Sequence 1897 BP; 435 A; 521 C; 532 G; 407 T; 2 other;
                                                                                                                                        of the invention.
                                                                                                                                                                                                                                                                                                  (CORI-) CORIXA CORP
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                                      GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
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                                                                                                                                                                                                                                                                               Stolk JA, Algate PA,
                                                                                      Similarity
                                                                                                                                                                                         Page 196; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                           gene;
                                                                                                                                                                                                                                                                                                                     ; 2000US-0713550.
; 2001US-0825294.
; 2001US-0970966.
                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     637
                                                                                       85.4%; Score 315;
100.0%; Pred. No.
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                                                                              Mismatches
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                                                                                        4.2e-158;
                                                                                                   DB 24;
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                                                                                                  Length 1897;
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RESULT 11
ABL40352
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     ABL40352;
The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. They are also useful for the isolated polynucleotides of the invention are useful for their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ovarian cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ovarian carcinoma 01034C/0591S consensus nucleotide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                              US2002004491-A1
                                                                                                                                               Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
                                                                                                                                                                                                                                                          (STOL/)
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15-AUG-2000;
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                                                                                                                        Claim 1a; Page 127-128; 131pp; English.
                                                                                                                                                                                    P-PSDB;
                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                      Xu J,
                                                                                                                                                                                                                                                                                                        14-NOV-2000;
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                                                                                                                                                                                                                                             XU J.
) STOLK J A.
) ALGATE P A.
) FLING S P.
                                                                                                                                                                                    ABB09417
                                                                                                                                                                                                                                                                                                        2000US-0561778.
2000US-0640173.
2000US-0656668.
2000US-0713550.
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                                                                                                                                                                                                                                                                                                                                                         99US-0394374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer therapy; vaccine; gene therapy; tumour; cancer;
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             25-APR-2001; 2001WO-US13788
                                             01-NOV-2001.
                                                                                                                                                                                                                                                                   Human; cardiovascular; tumourigenic disorder; aberrant angiogenesis; gene therapy; aberrant vascularisation; artherosclerosis; ovarian cancer; ischaemia/reperfusion injury; hypertension; arterial inflammation; psoriasis; endothelial cell disorder; diabetic retinopathy; restenosis; myocardial infarction; Grave's disease; G protein coupled receptor; GPCR 4941; vasottopic; hypotensive; antiinflammatory; cytostatic;
                                                                              WO200181634-A2
                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAD18690 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma O1034C/O591S consensus
                                                                                                                                                                                                                                                             antidiabetic; antipsoriatic;
                                                                                                                                                                                                                                                                                                                                                                                                                           18-DEC-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                          protein coupled receptor (GPCR) 4941 cDNA.
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                                                                                                          /product= "Human GPCR 4941"
/note= "The CDS is specifically claimed in
the specification"
                                                                                                                                                                            Location/Qualifiers 42..1403
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2528
                                                                                                                                                                                                                                                           leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                             SS
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                                                                                                                          calim 1 of
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AAF22400/c
ID AAF224
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AC AAF224
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DT 26-MAR
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KW Human;
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                                                                                                                                                                                                                                                                                                                                                                2037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  molecule (6 protein coupled receptor gene, GPCR 4941) associated with a cardiovascular or tumourigenic disorder. The method comprising contacting a sample containing a nucleic acid molecule with a hybridisation probe or amplification primers and detecting the presence. The invention is used in gene therapy. The method of the invention is used for identifying nucleic acids or polypeptides associated with a cardiovascular or tumourigenic disorder such as aberrant angiogenesis, aberrant vascularisation, artherosclerosis, or ovarian cancer, ischaemia/ reperfusion injury, hypertension, restenosis, arterial inflammation, endothelial cell disorders, diabetic retinopathy, psoriasis, myocardial infarction, Grave's disease and leukaemia. The methods can also detect mRNA or genomic DNA in a sample. The present sequence is 6 protein
                                                                                                                                         AAF22400 standard; cDNA;
                                                                                                                                                                                                                          1917 TGCCTCGGCCCTCA 1904
                                                                                                                                                                                                                                                                                                                                                                                                                                     2097
                                                                        26-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2217 AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG
                                                                                                                                                                                                                                                            331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2528 BP; 516 A; 766 C; 677 G; 567 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a method for identifying a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Fig 1; 118pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identifying nucleic acids for the diagnosis and treatment of cardiovascular and tumorigenic disorders, comprises identifying
                                                                                                                                                                                                                                                                                      151 AGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein coupled receptor (GPCR)-4941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-611743/70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-APR-2000;
09-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91 CTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGA 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31 AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                           TGCCTCGGCCCTCA 344
                                                                                                                                                                                                                                                                                                                                                          TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                                                                                                                                                                                                                                                                           TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGA
                                                                                                                                                                                                                                                                                                                                                                                                                             AGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTG
                                    secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         314;
                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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2000US-0635521.
                                  protein gene 28 SEQ ID NO:38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      85.1%;
100.0%;
                                                                                                                                         1953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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Pred. No. 1.4e-15
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Human; secreted protein; diagnosis; immunosuppressive; antiarthritic;

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                                                                                                                                                                                                                                                                      CC and polypeptides homologous to them. Human secreted protests have CC activities based on the tissues and cells the genes are expressed in. CC Examples of activities include: immunosuppressive; antiarthritic; CC antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; CC fungicide; and ophthalmological. The polypucleotides and proteins can be used to prevent, treat or ameliorate a medical condition in e.g. humans, CC mice, rabbits, goats, horses, cats, dogs, chickens or sheep. They are CC also used in diagnosing a pathological condition or susceptibility to a pathological condition. Disorders which are diagnosed or treated include alsorders e.g. neoplasms of the breast or liver, cardiovascular CC disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia, angiogenesis, nervous system disorders e.g. Alzheimer's CC disorders e.g. corneal infection. The polypeptides can also be used to add wound healing and epithelial cell proliferation, to prevent skin candiovative to increase or decrease storage capabilities. AAF22364 to AAF22372 and AAB63133 represent sequences used in the exemplification of the present invention.
                                                                                                                                                   Matches
 1585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; ophthalmological; gene therapy; autoimmune disease; neoplasm; rhumatoid arthritis; hyperproliferative disorder; cardiac arrest; cardiovascular disorder; cerebrovascular disorder; cerebral ischaemia; angiogenesis; nervous system disorder; Alzheimer's disease; infection; ocular disorder; corneal infection; wound healing; skin aging;
                                                                         1645
                                                                                                                                                                                                                       Sequence 1953 BP; 428 A; 540 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF22373 to AAF22421 encode the human secreted proteins given in AAB6313 to AAB63183. AAB63183 to AAB63231 represent more human secreted proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated nucleic acid molecule encoding a human secreted protein used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-APR-1999;
20-JAN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antirheumatic;
                                    90
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                                                                                                                                                                       Foca T
                                                                                                                                                                                                                                                              present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-647420/62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        additive;
                 GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
                                                                                                            CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
                                                                     CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 1586
                                                                                                                                                   283;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HUMAN GENOME SCI INC ROSEN C A.
                                                                                                                                                                     Similarity
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                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0128694.
2000US-0176931.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        preservative; ss.
                                                                                                                                                                   76.7%;
100.0%;
                                                                                                                                                 0;
                                                                                                                                                                 Score 283;
Pred. No.
                                                                                                                                                                                                                       524 G; 458 T; 3 other;
                                                                                                                                                 Mismatches
                                                                                                                                                                                      DB 21;
                                                                                                                                                                   3e-141;
                                                                                                                                                                                  Length
                                                                                                                                                 Indels
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1526
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AAF64188/c
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         CC Human secreted proteins AAB75506 - AAB75554 are encoded by polynucleotide CC sequences AAB764176 - AAB764224. The specification includes amino acid CC sequences AAB75555 - AAB75606 which represent fragments of the human CC secreted proteins, and protein sequences with which they share homology. CC The proteins and polynucleotides, their agonists and antagonists have CC activities dependent on the tissues and cells in which they are CC expressed, examples of these activities include, immunosuppressive; CC virucide; fungicide; antirheumatic; antiproliferative; cytostatic; cardiant; CC virucide; fungicide; opthalmalogical; and vulnerary. The proteins, CC virucide; fungicide; opthalmalogical; and vulnerary. The proteins of the polynucleotides, agonists and antagonists can be used to treat or detect cor diagnose various diseases and disorders including, autoimmune CC diseases e.g. rheumatoid arthritis, hyperproliferative disorders c.g. cardiac arrest, cerebrovascular disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders
                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 443-444; 542pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; opthalmalogical; vulnerary; autoimmune disease; cardiovascular disorder hyperproliferative disorders; cerebrovascular disorder; wound healing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1405
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                                                                                                                                                                                                                                                                                                                                                                                              Nucleic acid molecules encoding human secreted proteins, used in preventing, treating or ameliorating a disorder, e.g. Alzheimer's and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rosen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nervous system disorder; ocular disorder; skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted protein gene 13 SEQ ID NO:23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-APR-2001
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                                                                                                                                                                                                                                                                                                                                                                               Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ROSE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   additive; ss.
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                                                                                                                                                                                                                                                                                                                                                                             treating or ameliorat
diseases and cancers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunosuppressive; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chemotaxis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antirheumatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disorder;
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e.g. corneal infection.

The polypeptides can

also

used

to aid wound

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RESULT 15
AAF94818
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Best Local S
Matches 283
                                                                                                                                                                     10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  healing and epithelial cell proliferation, to prevent skin ageing due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotraxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities. Included in the invention are polynucleotide sequences AAF64167 - AAF64175 and peptide AAF7505 which are used in the isolation, identification and characterisation of the proteins of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1585
                                               Isolated polypeptides associated with ovarian carcinomas, nucleic acids that encode them, useful for the prevention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1465
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            Claim 5;
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                                                                                                                   Хu J,
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                                                                                                                                                                                                                                                                                            WO200118046-A2
                                                                                                                                                                                                                                                                                                                                               Human, ovarian
                                                                                                                                                                                                                                                                                                                                                                        Human
                                                                                                                                                                                                                                                                                                                                                                                                   23-MAY-2001
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                                                                                                                                            (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                     domo sapiens
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                                                                                        2001-211395/21.
                                                                                                                                                                                                                                                                                                                                                                        ovarian
                                                                                                                   Stolk JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
           Page 119;
                                    of ovarian cancers
                                                                                                                                                                     ; 2000US-0561778.
; 2000US-0640173.
; 2000US-0656668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                      cancer associated coding sequence
                                                                                                                                                                                                                                                                                                                                               cancer;
                                                                                                                                                                                                              99US-0394374
         189pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                      CDNA; 396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
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                                                                                                                                                                                                                                                                                                                                               vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 283; DB 22; 1
Pred. No. 5.3e-141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                              gene
                                                                                                                                                                                                                                                                                                                                             therapy;
                                                                                                                                                                                                                                                                                                                                               carcinoma; ss.
                                               carcinomas, and the prevention diagnosis
                                                                                                                                                                                                                                                                                                                                                                        SEQ
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                                                                                                                                                                                                                                                                                                                                                                        ID NO:
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RESULT 16
ABT03085
ID ABT03
XX
ABT03085
AC ABT03
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AC ABT03
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PR 03-
PR 11-
PR 03-
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Best Local Simi
Matches 281;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides a number of coding sequences and proteins, the over-expression of which is associated with overian carcinoma/cancer. These can be used in the diagnosis, treatment and prevention of ovarian cancer, optionally by gene therapy or in the form of a vaccine. The present sequence is an example of one of these sequences.
                                                                                                                                                          Novel ovarian cancer polypeptide and polynucleotide, useful fidetecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovar
                                                                                                                                                                                                                                                                                                                                                                                                                                 14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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                                                                                                                                                                                                                                                                                                                             Хu J,
                                                                                                                                                                                                                                                                                                                                                                            (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-NOV-2001; 2001WO-US45395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200239885-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human ovarian carcinoma associated coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABT03085 standard; cDNA; 396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCATCAGCTGCTGCAACACCCCTCTTTGTAACGGGCCAAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2002-500186/53.
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                                                                                                                                                                                                                                                                                                                           Stolk JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer;
                                                                                                                                                                                                                                                                                                                        Algate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         76.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ovarian carcinoma;
                                                                                                                                                                                                                                                                                                                             PΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                        Fling
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                                                                                                                                                                                                                                                                                                                             SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 22;
6e-140;
                                                                                                                                                                                                                                                                                                                             Molesh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy; immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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                                                                                                                                                                                                                                                                                                                             DA
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                                                                                                                                                                   ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                     cancer
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The present invention provides human over and coding sequences. The sequences can

ovarian

arian cancer be used in t

the associated

diagnosis and

Example 1;

Page 116; 197pp; English.

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RESULT 17
ABL48768
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Best Local :
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                                                                                                                                                                                         10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 396 BP; 90 A; 117 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                     (STOL/)
                                                                                                                                                                                                                                                                                                                                            Ovarian cancer; cancer therapy; vaccine; gene therapy; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                 Ovarian carcinoma sequence
                                                                                                                                                                                                                                                                                                                                                                                      18-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                          ABL48768
                                                                                                                                                                                                                                                                                                                                                                                                                               ABL48768 standard; cDNA;
                               Claim la; Page 41-42; 131pp; English.
                                                  Ovarian tumour polypeptide and prevention and/or treatment of
                                                                                   WPI; 2002-171027/22.
                                                                                                       Xu J,
                                                                                                                                                                                 14-NOV-2000;
                                                                                                                                                                                                                                               03-APR-2001; 2001US-0825294.
                                                                                                                                                                                                                                                                   10-JAN-2002.
                                                                                                                                                                                                                                                                                       US2002004491-A1
                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 210
The invention relates to ovarian tumour that may be utilised in cancer therapy,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    280
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention.
                                                                                                      Stolk JA,
                                                                                                                                    STOLK J A.
ALGATE P A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                            FLING S P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                        99US-0394374.
2000US-0561778.
2000US-0640173.
2000US-065668.
                                                                                                                                                                                 2000US-0713550
                                                                                                       Algate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        76.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                 396
                                                                                                        PΑ,
                                                                                                                                                                                                                                                                                                                                                                   isolate 21920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 281;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95 G; 92 T; 2 other;
                                                                                                       Fling
                                                   polynucleotide useful in diagnosis, cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6e-140
 polynucleotides and polypeptides for example in a vaccine or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      149
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RESULT 18
ABT03282
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ρy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ρ
THE PRESENTATION OF THE PR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The sequences given in records ABL48760-ABL48956 represent polynucleotides encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer is stimulating an immune response in a patient, and for treating a cancer is
                                                                                                                                                               14-NOV-2000;
03-APR-2001;
02-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 396 BP; 90 A; 117 C; 95 G; 92 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ovarian carcinoma proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABT03282 standard; cDNA; 1010 BP
                                                                                                                                                                                                                                                                              13-NOV-2001; 2001WO-US45395
                                                                                                                                                                                                                                                                                                                                                                                                 WO200239885-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-SEP-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABT03282;
                                                      Хu J,
                                                                                                                                                                                                                                                                                                                                          23-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         160
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                                                                                                            (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GCATCAGCTGCTACACCCCTCTTTGTAACGGGCCAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCATCAGCTGCTAACACCCCCTCTTTGTAACGGGCCAAGG 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       281;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ovarian cancer; ovarian carcinoma;
                                                      Stolk JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                               ; 2000US-0713550.
; 2001US-0825294.
; 2001US-0970966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    carcinoma
                                                         Algate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               76.2%; Score 281; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    associated coding sequence SEQ ID NO: 212.
                                                         PΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                         Fling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                         SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6e-140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 24;
                                                         Molesh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy;
                                                         DA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunotherapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     269
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'n
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WPI; 2002-500186/53

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XXX ACC XXX AC
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ABL40350
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                XXGCCCCX BX P P P P XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 271; Conserv
                                                                                                                                                                              01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                                                      (XUJJ/)
(STOL/)
(ALGA/)
(FLIN/)
       Хu Л,
                                                                                                                                                                                                                                                                                                                                                                                                                                  US2002004491-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ovarian carcinoma BF345141 nucleotide sequence.
                                                                                                                                                                                                                                                                                   10-SEP-1999;
                                                                                                                                                                                                                                                                                                                             03-APR-2001; 2001US-0825294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ovarian cancer; cancer therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABL40350 standard;
                                                                                                                                                                                                                                                                                                                                                                                 10-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1010 BP; 266 A; 344 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2; Page 195; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       271 CATCAGCTGCTGCAACACCCCTCTTTGTAAC 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                318
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91 CTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGA 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31 AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CATCAGCTGCTACACCCCCTCTTTGTAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGA
                                                    ) STOLK J A.
) ALGATE P A
) FLING S P.
     Stolk JA,
                                                                                                                               XU J.
                                                                                                                                                                           99US-0394374.
2000US-0561778.
2000US-0640173.
2000US-0656668.
2000US-0713550.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        73.4%;
ilarity 100.0%;
Conservative
                                                                           .
     Algate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cDNA;
     PΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1010
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b; Pred. No. 1.4
0; Mismatches
Fling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      254 G; 146 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  528
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 1010;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               coding sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90
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RESULT 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                               Primer specific for DNA encoding secretory/membrane protein SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma BF345141 nucleotide
                                                                 EP1067182-A2
                                                                                                    Synthetic
                                                                                                                                   Human; secretory protein; membrane protein; vaccine; gene therapy;
rheumatoid arthritis; diabetes; PCR primer; ss.
                                                                                                                                                                                                                          23-MAY-2001
                                                                                                                                                                                                                                                                                            AAF94044 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1010 BP; 266 A; 344 C; 254 G; 146 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim la; Page 126-127; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI;
                                                                                                                                                                                                                                                                                                                                                                                     498
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31 AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                    CATCAGCTGCTGCAACACCCCTCTTTGTAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CATCAGCTGCTACACCCCCTCTTTGTAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGA 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002-171027/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   271;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                73.4%; Score 271; DB 24
100.0%; Pred. No. 1.4e-1
                                                                                                                                diabetes; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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                                                                                                                                                                                                                                                                                                                                                                                  528
                                                                                                                                                                                                                                                                                                                                                                                                                   301
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       210
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    497
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07-JUL-2000; 2000EP-0114090

10-JAN-2001

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC which encode human scretcry or membrane protectins represented by

CC AAR83317 - AAR8419. Included in the invention are primers

CC AAR93917 - AAR94295 and AAR62232 - AAR62235 which are used to isolate the

CC CAAR93917 - AAR94295 and AAR62232 - AAR62235 which are used to isolate the

CC CDNA sequences of the invention. The invention also includes methods for

CC the production of antibodies directed against the proteins, and cDNA

CC sequences, which can be used in vaccines. The polynucleotide sequences

CC can be used in gene therapy. The polynucleotide sequences

CC diagnosis of diseases associated with inappropriate secretory

CC proteins they encode may be used in the prevention, treatment and

CC diagnosis of diseases associated with inappropriate secretory

CC sequences may also be used as DNA probes in diagnostic assays

CC (e.g. polymerase chain reactions (PCR)) to detect and quantitate the

CC presence of similar nucleic acid sequences in samples. They may also be

CC used to study the expression and function of secretory proteins/membrane

CC polypeptides and their role in metabolism. The polypeptides may be used

CC as antigens in the production of antibodies against them and in assays to

CC identify modulators (agonists and antagonists may also be used as diagnostic agents for detecting the presence of the

CC agents to down regulate expression and activity. The antibodies may also

CC be used as diagnostic agents for detecting the presence of the

CC polypeptides in samples (e.g. by enzyme linked immunosorbant assay

CC collection and diabetes.
 BXBXAXB
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                                                                                                                                                  RESULT 21
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Best Local :
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11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to nucleic acid sequences AAF93744 - AAF93916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ ID 478; 609pp + CD ROM; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acids encoding secretory proteins/membrane progene therapy or as candidate target molecules in drug
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 591 BP; 108 A; 198 C; 173 G; 109 T; 3 other;
Human ovarian antigen HVCAG04 cDNA,
                                       22-AUG-2002
                                                                                                               ABQ54231 standard; cDNA; 1608
                                                                                                                                                                                                                                                                                                                                                            383
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                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
                                                                                                                                                                                                                                                                                                                                                                                                                                      CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                                                                                                                                                                                                                         AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                                                                                            GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCA 543
                                                                                                                                                                                                                                               GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCA 250
                                                                                                                                                                                                                                                                                     AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Isogai T, Nishikawa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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2000JP-0118775.
2000JP-0183766.
                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     59.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 221; DB 22,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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                                                                                                                 ВP
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   SEQ ID NO:111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sugiyama T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 591;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            proteins, useful in rug development -
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                                                                                                                                                                                                                                                                                                                                                            442
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Human; ovarian antigen; ovary; ovarian; breast; cancer; tumour; ovarian cancer; breast cancer; tumour; reproductive system disorder; infertility; pregnancy disorder; anovulation; polycystic ovary syndroupcos; ovarian cyst; dysmenorrhoea; endocrine disorder; infection; inflammatory condition; immune disorder; blood disorder; cardiovascular disorder; respiratory disorder; neurological disorder; gastrointestinal disorder; urinary system disorder; draw screening; gene therapy; chromosome mapping; forensic analysis; antibody preparation; cytostatic; immunomodulatory; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                              gene;
                                                                                                                                                                                                                                                                                                                                                                                               antibody preparation; cytostatic; immunomodulatory; neuroprotective; antiinflammatory; gynaecological; reproductive; chromosome 2q21-22;
07-JUN-2001; 2001WO-US18569.
                                                                                         03-JAN-2002
                                                                                                                                                                                   WO200200677-A1
                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     syndrome;
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(HUMA-) HUMAN GENOME SCI INC

07-JUN-2000; 2000US-209467P

P-PSDB; ABP41154. 2002-147878/19

Isolated nucleic acid molecules encoding novel ovarian polypeptides, useful in the prevention, treatment and diagnosis of cancer (e.g. ovarian cancer), immune disorders, cardiovascular disorders and neurological diseases -

Claim 1; SEQ ID No 111; 2922pp; English.

CC ABP43228) and to cDNAs encoding them (ABQ54131-ABQ56305), and also CC encompasses polypeptides 90% identical and polynucleotides 95% identical content on the sequences of the invention. The invention additionally relates to recombinant vectors and host cells comprising human ovarian antigen CC polynucleotides, antibodies against human ovarian antigen CC iteating, prognosing or preventing various ovary and/or breast-related CC metastaris. Such conditions include ovarian corer and breast cancer, and cdisorders. Such conditions include ovarian cancer and breast cancer, and cdisorders. Such conditions include ovarian cancer and breast cancer, and cdisorders (e.g., infertility, disorders of pregnancy, anovulation, complexities, infections (e.g., chlamydia, HIV, toxoplasmosis, and toxic conspirations, immune disorders (e.g., congenital and acquired vaginitis), immune disorders (e.g., congenital and acquired conspiratory disorders (e.g., anaemia), cardiovascular disorders. CC inventions (e.g., anaemia), cardiovascular disorders. CC condulate ovarian antigen expression or activity. The polynucleotides may also be used in screening for compounds which modulate ovarian antigen expression or activity. The polynucleotides may be used as food additives or to prepare antibodies useful in disease diagnosis, drug targeting and phenotyping. The present sequence represents cDNA encoding a human ovarian antigen of the Note: The sequence data for this patent did not form specification, but was obtained in electronic format part of the printed directly from WIPO

Sequence 1608 BP; 402 A; 402 C; 417 G; 381 T; 6 other;

ftp.wipo.int/pub/published\_pct\_sequences.

Query Match
Best Local Similarity Matches Conservative 38.5%; Score 142; D 100.0%; Pred. No. 1. tive 0; Mismatches DB 24; . 2e-65; Length 1608; Indels 0 Gaps 0

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                                                                                                                                            Query Match
Best Local
                                                                                                                                 Matches 167;
                                                                                                                                                                                                                                                                                             Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                    14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
                                                                                                                                                                             Sequence 480 BP; 86 A; 154 C; 153 G; 87 T; 0 other;
                                                                                                                                                                                                      of the invention.
                                                                                                                                                                                                            The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                WPI; 2002-500186/53.
                                                                                                                                                                                                                                                                                                                                                                        Χu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-NOV-2001; 2001WO-US45395.
                                                                                                                                                                                                                                                              Claim 2; Page 196; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200239885-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; ovarian cancer; ovarian carcinoma; gene therapy; immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-SEP-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                              (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABT03283;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABT03283 standard;
                                     369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCAT
                                                                                   309
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                                                                                                       30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                 GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTGAGAAAG
                                            GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCTCAGAAAG 149
                                                                                 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTCTTTGTAACGGGCCAAGGCC 312
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAGCACACCC
ovarian carcinoma associated coding sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                     Stolk JA, Algate
                                                                                                                                            Similarity
                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cDNA;
                                                                                                                                       31.7%;
99.4%;
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                                                                                                                                          Score 117;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                     Fling SP,
                                                                                                                               Mismatches
                                                                                                                                          DB 24;
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                                                                                                                                                     Length 480;
                                                                                                                             Indels
           197
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GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG

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ABL40351
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                                                                  Matches
                                                                                Query Match
Best Local
                                                                                                                                                  useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma BE336607 nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                     The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (XUJJ/)
(STOL/)
(ALGA/)
                                                                                                                            Sequence 480 BP; 86 A; 154 C; 153 G; 87 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                    Claim la; Page 127; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Хu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (FLIN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ovarian cancer; cancer therapy; vaccine; gene therapy; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ovarian carcinoma BE336607 nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABL40351;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABL40351 standard; cDNA; 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         429 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCAT 476
                                   30
                                                                                Local Similarity
167;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stolk JA, Algate PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             XU J.
) STOLK J A.
) ALGATE P A
) FLING S P.
                                                                Conservative
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2000US-0640173.
2000US-0656668.
2000US-0713550.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2
                                                                               31.7%;
                                                                0;
                                                                           Score 117; DB 24; Length 480; Pred. No. 2.8e-52;
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                                                                Mismatches
                                                              Indels
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ABL81273
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                                                                                                                          The present invention describes a composition (I) comprising: carriers cc and immunostimulants; and a polypeptide (II) of a ovarian tumour cc polypeptide encoded by a polypucleotide (II) having a cDNA sequence cc (S1) from the 10912 nucleotide sequences as given in ABL/7023 to cc (S1) from the 10912 nucleotide sequences as given in ABL/7023 to cc (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to cc (S1) can be used for detecting ovarian cancer in a patient's biological cc sample preferably serum or ovarian tissue. The method comprises cc contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to patient, where the amount of polynucleotide hybridising to (IV) is cetected preferably by polymerase chain reaction (PCR). (I) comprising cc (III) and/or (II) is useful for stimulating and/or expanding T cells specific for an ovarian tumour protein comprising contacting T cells contacting to (IV) is useful for stimulating and/or expanding T cells contacting for (III) is useful in design and preparation of clls cultible library e.g., a tumour cDNA library using well known
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                                                           Query Match
Best Local
                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                  Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ovarian cancer; ovarian tumour; cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-MAY-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-DEC-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human ovarian cancer related cDNA clone SEQ ID NO:4251.
                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID 4251; 489pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-122075/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Algate PA, Harlocker SL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-MAY-2000; 2000US-207484P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAY-2001; 2001WO-US17756
                                                                                              Sequence 430 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   369 GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTGAGAAAG 428
AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCAT 476
                                                55;
                                                          Similarity
                                             14.9%;
ilarity 100.0%;
Conservative
                                                                                              92 A; 121 C; 108 G; 109 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jones
                                                0;
                                                          Score 55;
Pred. No.
                                                Mismatches
                                                          DB 24; 1
                                                 0
                                                                      Length 430;
                                                Indels
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                                                0
                                                Gaps
                                                0;
18-APR-2000;
19-MAY-2000;
07-JUN-2000;
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23-AUG-2000;
30-AUG-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
05-SEP-2000;
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14-AUG-2000;
18-AUG-2000;
22-AUG-2000;
22-AUG-2000;
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14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
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14-AUG-2000;
14-AUG-2000;
                                     08-SEP-2000;
08-SEP-2000;
08-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human reproductive system related antigen DNA SEQ ID NO: 10144
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07-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; reproductive system related antigen; reproductive system disorder;
                                                                                                                                                                                                                                                                                                                                                                                         26-JUL-2000;
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                                                                                                                                                                                                                                                                                 2000US-0225268.
2000US-0225270.
2000US-0225447.
                      2000US-0231242.
2000US-0231243.
2000US-0231244.
2000US-0231413.
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2000US-0215135.
2000US-0216647.
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2000US-0217487.
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2000US-0209467.
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2000US-0198123.
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2000US-0224519.
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2000US-0228924.
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2000US-0225758
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2000US-0230438
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2000US-0246474
2000US-0246475
2000US-0246478
2000US-0246524
2000US-0246524
2000US-0246525
2000US-0246527
2000US-0246528
2000US-0246528
2000US-0246610
2000US-0246611
2000US-0246611
2000US-0249207
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2000US-0249218
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2000US-0249216
2000US-0249265
2000US-0249265
2000US-0249297
2000US-0249297
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2000US-0241221.
2000US-0241785.
2000US-0241786.
2000US-0241787.
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2000US-0236802
2000US-0237037
2000US-0237039
2000US-0237039
2000US-0237030
2000US-023933
2000US-023933
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2000US-0241809.
2000US-0241826.
2000US-0244617.
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2000US-0234998.
2000US-0235484.
2000US-0235834.
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2000US-0236368.
2000US-0236369.
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2000US-0234274.
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2000US-0233065.
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2000US-0231968
RESULT 26
ABK62080/c
ID ABK620
XX ABK620
XX ABK620
XX ABK620
XX Human;
XW Human;
XW Human;
XW mening
KW mening
KW mening
KW melti;
KW autism
KW aleep
KW alutism
KW Sleep
KW Sleep
XX Homo:
XX Homo:
XX Homo:
XX Homo:
XX 11-AU
PFR 19-OC
XX
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Best Local :
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01-DEC-2000;

05-DEC-2000;

05-DEC-2000;

06-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;
              11-AUG-2000; 2000US-224642P
19-OCT-2000; 2000US-241779P
                                                                                                                                                                                 Human; ss; gene; secreted protein; cancer; autoimmune disease; arthritis; osteopprosis; Alzheimer's disease; Parkinson's disease; meningitis; encephalitis; neoplasia; trauma; ischaemia; infarction; mania; stroke; cardiovascular disease; atherosclerosis; sepsis; anaemia; rheumatoid arthritis; hypothyroidism; allergic response; liver failure; multiple sclerosis; haemorrhage; paranola; obsessive compulsive disorder; autism; panic disorder; learning disability; feeding disorder; sleep pattern disorder; balance; perception; Thi-dependent insulitis; adult respiratory distress syndrome; ARDS.
                                                              30-JUL-2001; 2001WO-US21124
                                                                                                                           WO200214358-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in the prevention and treatment of reproductive system disorders including cancer. The present sequence is a genomic sequence encuprotein of the invention.
                                                                                           21-FEB-2002
                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                           Human cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                  ABK62080 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1226 BP; 188 A; 438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides the protein and coding sequences of a number of human reproductive system related antigens. These can be used
                                                                                                                                                                                                                                                                                                                                                                     18-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                     ABK62080;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isolated nucleic acid molecule encoding a reproductive system antigen is used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-465570/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          317 AAAAGGGGAAGTTCTGCCT 335
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                      encoding novel secreted protein LP243(a).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000US-0251856
2000US-0251868
2000US-0251869
2000US-0251899
2000US-0251990
2000US-0254097
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2000US-0256719.
2000US-0251479.
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2000US-0250391.
2000US-0251030.
                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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Pred. No.
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14 - SEP - 2000
21 - SEP - 2000
21 - SEP - 2000
25 - SEP - 2000
25 - SEP - 2000
26 - SEP - 2000
27 - SEP - 2000
29 - SEP - 2000
20 - CCT - 2000
20 - CCT - 2000
20 - CCT - 2000
21 - CCT - 2000
22 - CCT - 2000
23 - CCT - 2000
24 - CCT - 2000
25 - SEP - 2000
26 - CCT - 2000
27 - CCT - 2000
28 - CCT - 2000
29 - CCT - 2000
20 - CCT - 2000
21 - NOV - 2000

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ABK62081/c
ID ABK62081 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel human secreted polypeptide having CC sequence 90% identical to the polypeptide sequences of LP105, LP061, CC LP224, LP240, LP239(a), LP243(a), LP243(b), LP253, LP218), LP251(a), CC LP252, LP239(b), LP235(a), LP255(a), LP244, LP186, LP251(b), LP255(b), CC proteins (included are the nucleic acids encoding the LP CC proteins (including complement, fragments encoding mature forms of the polypeptide or variant), a vector comprising the nucleic acid, a CC an anti-LP antibody, ant/agonists of LP and anti-LP-encoding mRNA CC ribozymes. The secreted protein or its agonist is useful in the CC disease (and in diagnosis), condition or disorder associated with CC aberrant levels of the secreted protein or disorder associated with CC arthritis, encephalitis, neoplasia, trauma, ischaemia and infarction, CC mania, stroke, cardiovascular disease, atherosclerosis, rhumatorial and infarction, CC arthritis, hypothyroidism, anaemia, sepsis, allergic responses, multiple sclerosis, liver failure, haemorrhages, paranola, obsessive compulsive CC disorder, autism, panic disorder, learning disabilities, ALS (amylotrophic lateral sclerosis) psychoses, disorders in feeding, cuseful for identifying compounds that bind to the secreted protein. The present sequence encodes a novel secreted protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                            Human; ss; gene; secreted protein; cancer; autoimmune disease; arthritis; osteoporosis; Alzheimer; s disease; Parkinson's disease; meningitis; encephalitis; neopilasia; trauma; ischaemia; infarction; mania; stroke; cardiovascular disease; atherosclerosis; sepsis; anaemia; rheumatoid arthritis; hypothyroidism; allergic response; liver failure; multiple sclerosis; haemorrhage; paranoia; obsessive compulsive disorder; autism; panic disorder; learning disability; feeding disorder; sleep pattern disorder; balance; perception; Th1-dependent insulitis;
                                 Homo sapiens
                                                                                                                                                                                                                                                        Human cDNA encoding novel secreted protein LP243(b)
                                                                                                                                                                                                                                                                                                18-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1811 AAAAGGGGAAGTTCTGCCT 1793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2080 BP; 303 A; 758 C; 676 G; 342 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 154-157;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel polypeptides and polynucleotides of secreted proteins useful for treating various diseases such as multiple sclerosis, cancer, autoimmune diseases, osteoporosis, Alzheimer's disease and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ELIL ) LILLY & CO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                     respiratory distress syndrome;
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19; Conser
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Pred. No. 6.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 2080;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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14-MAY-2002 ABL75913;

(first

entry)

multigene trait; plant breeding;

inheritance; characteristic; growth; development; disease resenvironmental adaptability; quality; yield; molecular marker;

corn tassel; gene;

corn tassel-derived polynucleotide;

cdps; hybrid breeding; CDPs;

resistance;

Corn tassel-derived polynucleotide (cdps) SEQ ID NO:5287.

밁

1953

AAAAGGGGAAGTTCTGCCT 1935

0

ABL75913 RESULT 28

ABL75913 standard;

cDNA; 237

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The invention relates to a novel human secreted polypeptide having CC sequence 90% identical to the polypeptide sequences of LP105, LP061, CC LP224, LP240, LP239(a), LP243(a), LP243(b), LP253, LP218), LP251(a), CC LP224, LP239(b), LP223(a), LP255(a), LP244, LP186, LP251(b), LP255(b), CC LP252, LP239(b), LP223(a), LP255(a), LP244, LP186, LP251(b), LP255(b), CC proteins (including complement, fragments encoding mature forms of the LP proteins (including complement, fragments encoding the LP proteins (including complement, fragments encoding mature forms of the proteins (including complement, fragments encoding mature forms of the protein solvent for the protein of the protein and comprising the vector, the preparation of the protein and compressive for the protein encoding maked compounds to the secreted protein or its agonist is useful in the compressive of the secreted protein e.g. cancer, autoimmune diseases, are protein e.g. cancer, autoimmune diseases, compounds the vector of disease, parkinson's disease, carthitis, osteoporosis, Alzheimer's disease, Parkinson's disease, carthitis, stroke, cardiovascular disease, atherosclerosis, rheumatoid compounds troke, cardiovascular disease, atherosclerosis, rheumatoid compounds that protein e.g. cancer in feeding, campounds that bind to the secreted protein. The compounds that bind to the secreted protein. The present sequence encodes a novel secreted protein of the invention.
                                      Best Loc
Matches
                                                                           Query Match
                                                                                                          Sequence 2222 BP; 317 A; 827 C; 733 G; 345 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 160-163; 235pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           autoimmune diseases, osteoporosis, Alzheimer's disease and Parkinson's
disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel polypeptides and polynucleotides of secreted proteins useful for treating various diseases such as multiple sclerosis, cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P-PSDB; AAU91329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-AUG-2000; 2000US-224642P
19-OCT-2000; 2000US-241779P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-FEB-2002
317 AAAAGGGGAAGTTCTGCCT 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ELIL ) LILLY & CO
                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002-304057/34.
                                        19;
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                                                        Similarity
                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Micanovic
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                                                      5.1%; Score 19; 100.0%; Pred. No.
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                                    0;
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                                    Mismatches
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                                                    DB 24;
6.8;
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                                    0
                                                                       Length 2222;
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                                    Indels
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            with growth and development, disease resistance, environmental adaptability, quality and yield, and as molecular markers for studying inheritance of multigene traits in a plant breeding program. (I) can be used to produce a tassel-specific profile of gene transcription, a transcript image, to clone regulatory elements for use in transformation vectors, to express a polypeptide, to identify, isolate or extend identical or related corn tassel nucleic acid sequences from DNA ilbraries, in nucleic acid hybridisation or amplification technologies, as query sequences to determine homology of known sequences, as probe for use in Southern or Northern hybridisation, and to identify the presence of and/or to determine the degree of similarity between two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encode corn tassel-dérived polypeptides (CDPs). The cdps sequences (I) can be used for determining altered gene expression, to recover regulatory elements and to follow inheritance of desirable characteristics through hybrid breeding programs. (I) are also useful in the evaluation, and alteration of desired characteristics associated in the evaluation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel purified corn tassel-derived polynucleotide useful for determining altered gene expression, to recover regulatory elemento follow inheritance of desirable characteristics through hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-163647/21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence describes a purified corn tassel-derived polynucleotide sequence (cdps) comprising a nucleic acid sequence selected from those given in ABL70627 to ABL76833. The cdps sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID 5287; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      breeding programs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lalgudi RV, Ito LY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-APR-1998;
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                                                 Corn; corn tassel-derived polynucleotide; cdps; hybrid breeding; CDPs; inheritance; characteristic; growth; development; disease resistance; environmental adaptability; quality; yield; molecular marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 237
                                    multigene trait;
                                                                                                                      Corn tassel-derived polynucleotide (cdps) SEQ ID NO:5709.
                                                                                                                                                            14-MAY-2002
                                                                                                                                                                                              ABL76335;
                                                                                                                                                                                                                                 ABL76335 standard; cDNA; 256
 Zea mays
                                                                                                                                                                                                                                                                                                                                                        192 GTGCATCATCAGCGGCCT 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 or more) nucleic acid sequences.
                                                                                                                                                                                                                                                                                                                       44 GTGCATCATCAGCGGCCT 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ITO L Y.
SHERMAN B K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LALGUDI R V.
                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 60 A; 64 C; 66 G; 46 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98US-082567P
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                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                               4.98;
                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                             Score 18;
                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                              DB 24; Length 237; 22;
                                                                                                                                                                                                                                                                                                                                                                                             0;
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ABL75647
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            with growth and development, disease resistance, environmental adaptability, quality and yield, and as molecular markers for studying inheritance of multigene traits in a plant breeding program. (I) can be used to produce a tassel-specific profile of gene transcription, a transcript image, to clone regulatory elements for use in transformation vectors, to express a polypeptide, to identify, isolate or extend identical or related corn tassel nucleic acid sequences from DNA ilbraries, in nucleic acid hybridisation or amplification technologies, as guery sequences to determine homology of known sequences, as probe for use in Southern or Northern hybridisation, and to identify the presence of and/or to determine the degree of similarity between two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence describes a purified corn tassel-derived polynucleotide sequence (cdps) comprising a nucleic acid sequence selected from those given in ABL70627 to ABL70833. The cdps sequences encode corn tassel-derived polypeptides (CDPs). The cdps sequences (I) can be used for determining altered gene expression, to recover regulatory elements and to follow inheritance of desirable characteristics through hybrid breeding programs. (I) are also useful in the evaluation, and alteration of desirable characteristics through hybrid breeding programs.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-163647/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (LALG/) LALGUDI R V.
(ITOL/) ITO L Y.
(SHER/) SHERMAN B K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      determining altered gene expression, to recover regulatory to follow inheritance of desirable characteristics through
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lalgudi RV, Ito LY,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel purified corn tassel-derived polynucleotide useful for
                                              Corn; corn tassel-derived polynucleotide; cdps; hybrid breeding; CDPs inheritance; characteristic; growth; development; disease resistance; environmental adaptability; quality; yield; molecular marker; multigene trait; plant breeding; corn tassel; gene; ss.
                                                                                                                                     Corn tassel-derived polynucleotide (cdps) SEQ ID NO:5021.
                                                                                                                                                                          14-MAY-2002
                                                                                                                                                                                                             ABL75647
                                                                                                                                                                                                                                              ABL75647 standard; cDNA; 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 256 BP; 72 A; 60 C;
                                                                                                                                                                                                                                                                                                                                                                        192 GTGCATCATCAGCGGCCT 209
                                                                                                                                                                                                                                                                                                                                       77
                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   more) nucleic acid sequences.
                                                                                                                                                                                                                                                                                                                                                                                                           . Similarity
18; Conser
                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                        Score 18; DB 2; Pred. No. 22; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  78 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  42 T; 4 other;
                                                                                                       cdps; hybrid breeding; CDPs;
                                                                                                                                                                                                                                                                                                                                                                                                                                              24; Length 256;
                                                                                                                                                                                                                                                                                                                                                                                                             0;
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hybrid
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RESULT 31
AAK66339/c
ID AAK663
XX
AC AAK663
XX
DT 06-NOV
XX
DT UMMAN
DE HUMMAN
KW HUMMAN;
KW CYTOST
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OS HOMO S
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PN WO2001
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PD 09-AUG
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                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                     characteristics through hybrid breeding programs. (I) are also useful in the evaluation, and alteration of desired characteristics associated with growth and development, disease resistance, environmental adaptability, quality and yield, and as molecular markers for studying inheritance of multigene traits in a plant breeding program. (I) can be used to produce a tassel-specific profile of gene transcription, a transcript image, to clone regulatory elements for use in transformation vectors, to express a polypeptide, to identify, isolate or extend identical or related corn tassel nucleic acid sequences from DNA libraries, in nucleic acid hybridisation or amplification technologies, as query sequences to determine homology of known sequences, as probe for use in Southern or Northern hybridisation, and to identify the presence of and/or to determine the degree of similarity between two
                            WO200157182-A2
                                                                              Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
cytostatic; gene therapy; vaccine; metastasis; ds.
                                                                                                                      Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21151
                                                                                                                                                    06-NOV-2001
                                                                                                                                                                               AAK66339;
                                                                                                                                                                                                        AAK66339 standard;
                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      encode corn tassel-derived polypeptides (CDPs). The cdps sequences can be used for determining altered gene expression, to recover regulatory elements and to follow inheritance of desirable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polynucleotide sequence (cdps) selected from those given in AF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID 5021; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel purified corn tassel-derived polynucleotide useful for determining altered gene expression, to recover regulatory elements and to follow inheritance of desirable characteristics through hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-163647/21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (LALG/) LALGUDI R V.
(ITOL/) ITO L Y.
(SHER/) SHERMAN B K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-APR-1998;
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                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-APR-1999;
                                                                                                                                                                                                                                                                                                    192 GTGCATCATCAGCGGCCT 209
                                                                                                                                                                                                                                                                           Local Similarity
les 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                present sequence describes a purified corn tassel-derived ynucleotide sequence (cdps) comprising a nucleic acid sequence ected from those given in ABL70627 to ABL76833. The cdps sequences
                                                                                                                                                                                                                                                                                                                                                                                                                more) nucleic acid sequences.
                                                                                                                                                                                                                                                                                                                                                                                   286 BP;
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                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                      69
                                                                                                                                                                                                        DNA; 484
                                                                                                                                                                                                                                                                                                                                                                                    A; 74 C;
                                                                                                                                                                                                                                                                                                                                           4.9%; Score 18;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sherman
                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                    83 G;
                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                    52 T; 8 other;
                                                                                                                                                                                                                                                                                                                                           DB 24; Length 286; 23;
                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                              0;
14 - AUG - 2000
18 - AUG - 2000
22 - AUG - 2000
22 - AUG - 2000
23 - AUG - 2000
23 - AUG - 2000
20 - SEP - 2000
21 - SEP - 2000
20 - SEP - 2000
21 - SEP - 2000
22 - SEP - 2000
23 - SEP - 2000
24 - SEP - 2000
26 - SEP - 2000
27 - SEP - 2000
28 - SEP - 2000
29 - SEP - 2000
20 - SEP - 2000
20 - SEP - 2000
21 - SEP - 2000
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14-SEP-2000
14-SEP-2000
14-SEP-2000
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25-SEP-2000
25-SEP-2000
27-SEP-2000
27-SEP-2000
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07-JUL-2000;
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19-MAY-2000;
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                                                                                                                                                                                                                                                                                               2000US-0230438.
2000US-0231242.
2000US-0231243.
2000US-0231244.
2000US-0231244.
                                                              2000US-0234274.
2000US-0234997.
2000US-0234998.
2000US-0235484.
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2000US-0229509.
2000US-0229513.
2000US-0230437.
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                                                                                                                                  2000US-0233064
2000US-0233065
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2000US-0225270
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2000US-0225267
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and

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13-OCT-2000
20-OCT-2000
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29-SEP-2000;
02-OCT-2000;
02-OCT-2000;
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01-DEC-2000
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08-DEC-2000
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17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
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02-OCT-2000;
WPI; 2001-483426/52
                                         Rosen CA,
                                                                                   HUMAN
                                         Barash SC,
                                                                                                                                                                                                         2000US -0236369
2000US -0236370
2000US -0236802
2000US -0237037
2000US -0237037
2000US -0237039
2000US -02349960
2000US -0241786
2000US -0241786
2000US -0241809
2000US -02418126
2000US -0246174
2000US -0246474
2000US -0246474
2000US -0246477
2000US -0246477
2000US -0246477
2000US -0246477
2000US -0246478
2000US -0246477
2000US -0246673
2000US -0246673
2000US -0246524
2000US -0246525
2000US -0246526
2000US -0249211
                                                                                                                          2000US-0251989.
2000US-0251990.
2000US-0254097.
2000US-0259678.
                                                                                   GENOME
                                                                                   SCI INC
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AAK66340/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) can mino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) croteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome consumed the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) concerns and caids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotietics may be used to produce the secreted (I), by inserting concerns and cancer metastases of haematopoietic-related diseases, especially concerns and cancer metastases of haematopoietic antigen genomic conserns record for the present invention. AAK54942 to AAK87690 and AAM82169 concerns read and the area in the area of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 18
31 - JAN-2000;
04 - FEB - 2000;
24 - FEB - 2000;
10 - MAR - 2000;
11 - MAR - 2000;
11 - MAY - 2000;
11 - MAY - 2000;
11 - MAY - 2000;
27 - JUN - 2000;
28 - JUN - 2000;
27 - JUL - 2000;
11 - JUL - 2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis -  \frac{1}{2} 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAK66340 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200157182-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; immune; haematopoietic; immune/haematopoietic antigen; cancer; cytostatic; gene therapy; vaccine; metastasis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21152.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAK66340;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 484 BP; 124 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-JAN-2001; 2001WO-US01354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              143 CAGAAAGAAGTGATGGAG
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2000US-0186350
2000US-019874
2000US-0199076
2000US-0198123
2000US-0209467
2000US-0214886
2000US-0214886
2000US-0214647
2000US-0216647
2000US-0216480
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA; 484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 22; Length 484; 23;
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14-AUG-2000; 14-AUG-2000; 14-AUG-2000;

26-JUL-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000;

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08-NOV-2000;
17-NOV-2000;
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17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
       AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis an treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, alignose and treat immune/haematopoietic-related diseases, especially anxiety.
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                                                                                                                                                                                                                                                      Nucleic acids encoding useful for preventing, metastasis -
                                                                                                                                                                                                                      Disclosure;
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2000US-0246477.
2000US-0246524.
2000US-0246524.
2000US-0246526.
2000US-0246526.
2000US-0246527.
2000US-0246610.
2000US-0246611.
2000US-0246611.
2000US-0246613.
2000US-0246613.
2000US-0249207.
2000US-0249207.
2000US-0249210.
2000US-0249211.
2000US-0249211.
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2000US-0249211.
2000US-0249212.
2000US-0249213.
2000US-0249214.
2000US-0249214.
2000US-0249215.
2000US-0249214.
2000US-0249214.
2000US-0249215.
2000US-0249214.
2000US-0249244.
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2000US-0251391
2000US-0251030
2000US-02517198
2000US-0251479
2000US-0251479
2000US-0251856
2000US-0251868
2000US-0251869
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2000US-0251899
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2000US-0251990
2000US-0251997
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 cancer
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diagnosing
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of.
 haematopoietic-derived
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es, especially cells. AAK64703

polypeptides, and

and

2000US-022064 2000US-0224518 2000US-022513 2000US-022526 2000US-022526 2000US-022526 2000US-022575 2000US-022934 2000US-022934 2000US-022934 2000US-023124 2000US-023124 2000US-023124 2000US-023124 2000US-023126 2000US-024126 2000US-024126

14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
18-AUG-2000;
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22-AUG-2000;
23-AUG-2000;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                 AAH52304 to AAH53970 represent nucleic acids (I) encoding polypeptides (II), given in AAGB1454 to AAGB3120, from Staphylococcus epidermidis. (I) and (II) can have antibacterial activity and therefore can be used in vaccination. The nucleic acids (I) may be used to produce the S. epidermidis polypeptides (II) via the production of vectors containing them which are used to produce hosts cells which express the polypeptides. The polypeptides (II) (and/or nucleic acids) may then be used to vaccinate subjects and to raise antibodies against the bacteria. The polypeptides may also be used to assay for other inhibitors of their activity and therefore identify compounds that may be used for the treatment of S. epidermidis infections, e.g. endocarditis. AAH53971 to AAH55090 represent specifically claimed S. epidermidis genomic DNA polynucleotide sequences from the present invention. AAH55091 to
                                  however the sequence listing only goes up to SEQ ID NO:4454 so even though sequences are given in the disclosure for SEQ ID NO:4465 to 44 no sequences are present for SEQ ID NO:4465 to 44
                                                                          AAH55098 represent oligonucleotide sequences and primers which are used in the exemplification of the present invention.

N.B. The present invention specifically claims all the polynucleotide sequences given in the sequence listing of the present specification,
                              no sequences are present for SEQ ID NO:4455 to 4464.
                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 8; Page 144; 2188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P-PSDB; AAG81646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kimmerly WJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GLAX ) GLAXO GROUP LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vaccination; endocarditis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Staphylococcus epidermidis SR1 strain; infection; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH52496
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to AAK87694 represent human immune/haematopoietic antigen genomic sequences from the present invention. AAK54942 to AAK54950 and AAM82169 represent sequences used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   457 CAGAAAGAAGTGATGGAG 440
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          for vaccinating
                                                                                                                                                                                                                                                                                                                                                                                                                                                acids encoding polypeptides from Staphylococcus epidermidis, for vaccinating against infections, e.g. endocarditis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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23;
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                                             NO:4465 to 4472
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Sequence 1152 BP; 429 A; 169 C;

241 G;

313 T; 0 other;

Query Match

Sequence 1542

BP; 416 A; 434 C; 433 G;

259 T; 0 other;

Best Local Similarity

100.0%;

Score 18; Pred. No.

DB 23; 23;

Length 1542;

4.98;

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RESULT 34
AAS83729/c
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                                                                                                                                                                          for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits to assess biodiversity
and to produce other types of data and products dependent on DNA and amino acid sequences. AAS64197-AAS94564 represent novel human diagnostic coding sequences of the invention. Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at ftp. wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to isolated polynucleotide (I) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutatio responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Drmanac RT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAR-2000; 2000US-0540217.
23-AUG-2000; 2000US-0649167.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; chromosome mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA encoding novel human diagnostic protein #19533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-FEB-2002 (first entry)
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hes 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001-639362/73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID No 19533; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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100.0%;
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maging; diagnostic; genetic (
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Pred. No.
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23;
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                                                                                                                                                                  ABN90538 to ABN93374 represent Staphylococcus epidermidis open reading frame (ORF) nucleic acid sequences which encode the amino acid sequences given in ABp35124 to ABp37960. The S. epidermidis sequences have antibacterial activity and can be used in gene therapy. The sequences can also be used in the diagnosis and treatment of bacterial infections, particularly S. epidermidis infections. The sequences can be used to screen for compounds able to interfere with the S. epidermidis life cycle or inhibit S. epidermidis infection.

N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from the
                                                                                                                                                                                                                                                                                                                                                                   14-AUG-1997;
08-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Staphylococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                              antibacterial;
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                                                                                                                                           Sequence 1587
                                                                                                                                                                                                                                                               Disclosure; SEQ ID 1425;
                                                                                                                                                                                                                                                                               Novel isolated nucleic acid encoding a polypeptide, useful for diagnosing and
                                                                                                                                                                                                                                                                                                                                   Doucette-Stamm
 24 - JUL - 2002
                                  ABN92620
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                                                                                                                                                                                                                                                                                                                                                  (GENO-) GENOME
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                                                                                                                  Local Similarity
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                                  standard;
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(first entry)
                                                                                                                                             BP; 605 A; 231
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97US-064964P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy;
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                                                                                                                                             325 G; 426 T; 0 other;
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                                                                                                                                                                                                                                                                                Staphylococcus epidermidis treating bacterial infections
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23;
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                      Human; open reading frame; ORFX; detection; cytostatic; hepatotropic; vulnerary; antipsoriatic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; osteopathic; antiarthritic; immunosuppressant; cardiant; immunostimulant; thrombolytic; coagulant; vasotropic; antidiabetic; hypotensive; dermatological; immunosuppressive; antiinflammatory; antiviral; antibacterial; antifungal; antitheumatic; antithyoid; antianaemic; gene therapy; cancer; proliferative disorder; hypertension; neurodegenerative disease; diabetes mellitus; hypothyoidism; SCID; ALDS; coholesterol ester storage; systemic lupus erythematosus; infection; cholesterol ester storage; systemic lupus erythematosus; infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cycle or inhibit S. epidermidis infection.

N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel isolated nucleic acid encoding a Staphylococcus epidermidis polypeptide, useful for diagnosing and treating bacterial infections
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     severe
                                                                                                                                                                                                                                                                                                                            Human ORFX ORF538 polynucleotide sequence SEQ ID NO:1075.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC74983 standard;
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18; Conserv
combined immunodeficiency; malaria;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 24;
        autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 1596;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bacterial infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO: 2083.
        disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
           asthma
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AAA55968/
ID AAA5
                                                                                                                                                                                                                                                                                                                                                                                                                SX FX BX S
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                                                                                                                                                                                   RESULT 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                             which represent the human OFFX open reading frames I to 3161. The OFFX CC sequences have activities such as: cytostatic; hepatotropic; vulnerary; cantipsoriatic; antiparkinsonian; nocytostatic; hepatotropic; vulnerary; costeopathic; anticonvulsant; antiarthritic; inmunosuppressant; immunosuppressant; immunosuppressant; immunosuppressant; immunosuppressant; immunosuppressant; cantiinflammatory; antibacterial; antiviral; antifungal; antirheumatic; antiinflammatory; antibacterial; antiviral; antifungal; antitheumatic; antithyroid; and antianaemic. The sequences can be used for determining the presence of or predasposition to, or preventing or treating character. The nucleic acids can be used to express ORFX proteins in gene therapy coefficiency. The proteins and nucleic acids may be used to treat cancers, proliferative disorders, neurodegenerative disorders, osteoarthritis, graft vs host disease, cardiovascular disease, diabetes mellitus, bypertension, hypothyroidism, cholesterol ester storage, systemic lupus crythematosus, severe combined immunodeficiency (SCID), AIDS, viral, bacterial or fungal infection, malaria, autoimmune disorders, asthma, concturnal haemoglobinuria, antiinflammatory disease; to enhance
                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                Matches
               Murine G713 encoding cDNA SEQ ID NO:6
                                                        05-SEP-2000
                                                                                                 AAA55968;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAC74446 to AAC77606 encode the proteins given in AAB40237 to AAB43397, which represent the human ORFX open reading frames 1 to 3161. The ORFX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel nucleic acids and peptides derived from open reading frame {\sf X}, useful for treating e.g. cancers, proliferative disorders,
                                                                                                                                      AAA55968 standard; cDNA; 1791
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1633 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 5; Page 1013-1014; 5507pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-602362/57.
P-PSDB; AAB40774.
                                                                                                                                                                                                                                                                                                                                                                                                                                                coagulation; to inhibit thrombosis; and as a contraceptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neurodegenerative disorders and cardiovascular disease -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-APR-1999; 99US-0127728.
30-MAR-2000; 2000US-0540763.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAR-1999;
02-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-MAR-2000; 2000WO-US08621.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200058473-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             allergy; aplastic anaemia; nocturnal haemoglobinuria; burn; wound;
bone damage; cartilage damage; antiinflammatory disease; coagulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                             315 ACCAGTCCTTCTGCTCCC 332
                                                                                                                                                                                                                                                                                    231 ACCAGTCCTTCTGCTCCC 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                thrombosis; contraceptive; ss.
                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                             Conservative
                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0127607.
99US-0127636.
                                                                                                                                                                                                                                                                                                                                                                                                         335 A; 515 C; 480 G; 303 T;
                                                                                                                                                                                                                                                                                                                   4.9%; Suc
100.0%; Pr
.... 0;
                                                                                                                                                                                                                                                                                                                                               Score 18; DB 21; Length 1633; Pred. No. 23;
                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                           other;
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                                                                                                                                  AAS91414/c
                                                                                                                                                     RESULT
                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes an isolated, purified or recombinant CC polynucleotide (PN) (I) comprising a contiguous span of 8 to 50 CC nucleotides, where the span includes a G713 or chromosome 13q31-q33 CC related biallelic marker. (I) has neuroleptic activity and can be used as a G713 gene expression inhibitor. (I) can be used genotyping to cestimate the frequency of an allele of a G713 or chromosome 13q31-q33 CC related biallelic marker in a population, and of a haplotype for a set CC of biallelic markers in a population. (I) is also useful in detecting an association between a haplotype and a trait. The frequency is used CC for detecting an association between a genotype and a trait being CC schizophrenia. The genotype is used to determine whether an individual cis at risk of developing schizophrenia. (I) can also be used as a CC medicament against several disorders preferably brain, psychiatric CC identification of risk of developing schizophrenia is possible, which CC would enable early and/or prophylactic treatment. Ana55964 to Ana55966 CC collabolic markers A12 to AA455030 represent human chromosome 13q31-q33 locus biallelic markers A12 to AA45; AAA5596 to AAA55991, and AAA55031 and CC AAY90963; AAA55962 to AAA55030 represent human chromosome 13q31-q33 locus invention.
                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                 Best
DNA encoding novel human diagnostic protein #27218
                                      13-FEB-2002
                                                                          AAS91414;
                                                                                                               AAS91414 standard; cDNA; 2253
                                                                                                                                                                                                                                                                                                                                                    Sequence 1791 BP; 389 A; 531 C; 508 G; 363 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel polynucleotide of human G713 gene useful for diagnosis and prophylactic treatment of brain, psychiatric disorders like schizophrenia and bipolar disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; chromosome 13; G713; chromosome 13q31-q33; schizophrenia; biallelic marker; polymorphism; central nervous disease; detecti neuroleptic; G713 gene expression inhibitor; genotyping; brain disorder; psychiatric disorder; bipolar disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 234-236; 271pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-OCT-1998;
30-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus.
                                                                                                                                                                                                         695
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                                                                                                                                                                                                                                         84 ACGACTGCTCCTCCCCC 101
                                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                                                                                                                                              18;
                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                            Conservative
                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bougueleret L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0103955.
98US-0106457.
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                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                    4.98;
                                                                                                                                                                                                       678
                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                              Score 18;
                                                                                                                                                                                                                                                                                               Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chumakov I,
                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                             DB
23;
                                                                                                                                                                                                                                                                                                              21; Length 1791;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cohen
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AAH54960/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polypeptide (II) sequences. (I) is useful as hybridisation probes, 2C polyperase chain reaction (PCR) primers, oligomers, and for chromosome 2C and gene mapping, and in recombinant production of (II). The 2C and gene mapping, and in recombinant production of (II). The 2C constitution of interest the production of the production of constant production partners are useful in medical constant protein expression or biological activity. (II) and (II) are useful for treating constant protein expression or biological activity. The polypeptide and polypuclacities sequences have applications in constant product expression or biological activity. The polypeptide and polypuclacities sequences have applications in constant product of the resist of assess biodiversity and to produce other types of data and products dependent on DNA and constant product sequences of the invention.

Constant product of the product of the printed constant printed printed constant printed printed constant printed printed constant prin
                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local :
Staphylococcus epidermidis SR1 strain; infection; vaccination; endocarditis; {\tt ds.}
                                                                 S. epidermidis
                                                                                                               03-SEP-2001
                                                                                                                                                            AAH54960
                                                                                                                                                                                                AAH54960 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2253 BP; 541 A; 612 C; 527 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAR-2000;
23-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-MAR-2001; 2001WO-US08631
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                             739
                                                                                                                                                                                                                                                                                                                                                    91 CTCCTCCCCGAGTTCAT 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                           CTCCTCCCCCGAGTTCAT 722
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2001-639362/73
                                                                                                                                                                                                                                                                                                                                                                                                 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABG27227.
                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Liu C,
                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-0540217.
2000US-0649167.
                                                               genomic polynucleotide sequence SEQ ID NO:4324.
                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27218; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          4.98;
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                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                 Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polynucleotide (I) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 573 T;
                                                                                                                                                                                                                                                                                                                                                                                                                 DB
23;
                                                                                                                                                                                                                                                                                                                                                                                                                                      23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0 other;
                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 2253;
                    diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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RESULT 4:
AAI59883
ID AAI!
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH52304 to AAH53970 represent nucleic acids (I) encoding polypeptides (II), given in AA681454 to AA683120, from Staphylococcus epidermidis. (I) and (II) can have antibacterial activity and therefore can be used in vaccination. The nucleic acids (I) may be used to produce the sequences in vaccination. The nucleic acids (I) may be used to produce the containing them which are used to produce hosts cells which express the polypeptides. The polypeptides (II) (and/or nucleic acids) may then be used to vaccinate subjects and to raise antibodies against the bacteria activity and therefore identify compounds that may be used for the rectivity and therefore identify compounds that may be used for the treatment of S. epidermidis infections, e.g. endocarditis. AAH53971 to AAH55090 represent specifically claimed S. epidermidis genomic DNA polynucleotide sequences from the present invention. AAH55091 to AAH55098 represent oligonucleotide sequences and primers which are used in the exemplification of the present invention. AAH55091 to AAH55098 represent livention specifically claims all the polynucleotide sequences and primers which are used in the exemplification of the present invention. AAH55091 to be sequences given in the sequence listing of the present specification, though sequences are given in the disclosure for SEQ ID NO:4454 so even though sequences are present for SEQ ID NO:4455 to 4461.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                Human: nootropic; immunosuppressant; cytostatic; gene therapy; cancer; peripheral nervous system; neuropathy; central nervous system; CNS; Alzhalmer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;
                                                                                                                                                                                                                                                                                                                                                                     1118
                                                                   chemokinetic;
                                                                                                                                                                      Human polynucleotide SEQ ID NO 3872.
                                                                                                                                                                                                                                              AAI59883;
                                                                                                                                                                                                                                                                              AAI59883 standard; cDNA; 3556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 3281 BP; 1226 A; 464 C; 636 G;
                                               leukaemia; ss
                                                                                                                                                                                                           22-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-316495/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-NOV-2000; 2000WO-US30782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200134809-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Staphylococcus epidermidis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GLAX ) GLAXO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                    269 TGCATCAGCTGCTGCAAC 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                   TGCATCAGCTGCTGCAAC 1101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               acids encoding for vaccinating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 2062-2063; 2188pp; English.
                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                thrombolytic; drug screening; arthritis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GROUP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polypeptides from Staphylococcus epidermidis, against infections, e.g. endocarditis -
                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 18;
Pred. No.
                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          955 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22;
                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 3281;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Homo sapiens

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RESULT 42
AAH54943/c
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                                                                                                                                                                          В
                                                                                                                                                                                                                         Query Match
Best Local 9
                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-JAN-2000;
25-APR-2000;
09-JUL-2000;
19-JUL-2000;
03-AUG-2000;
                                                                                                                                                                                                                                                                                                  The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Immune system suppression, Activinyinhibin activity, chemotactic/chemokinetic activity, hammostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200153312-A1
                                                                                                                                                                                                                                                                                          assays for receptor activity, arthritis C.N.S disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 3872; 10078pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel nucleic acids and such as central nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-DEC-2000;
WO200134809-A2
                  Staphylococcus epidermidis
                                             Staphylococcus epidermidis SR1
                                                                S. epidermidis
                                                                                     03-SEP-2001
                                                                                                                           AAH54943 standard; DNA; 3633
                                                                                                                                                                                                                                                      Sequence 3556 BP; 776 A; 1071 C; 1057 G; 652 T; 0 other;
                                                                                                                                                                                                                                                                        specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zhao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-SEP-2000;
19-OCT-2000;
                                     vaccination;
                                                                                                        AAH54943;
                                                                                                                                                                                                                                                                                  Note: The sequence data for this patent did not form part of the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HYSE-)
                                                                                                                                                                          629 ACCAGTCCTTCTGCTCCC 646
                                                                                                                                                                                             231 ACCAGTCCTTCTGCTCCC 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Š, Ž
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001-442253/47.
DB; AAM40727.
                                                                                                                                                                                                                l Similarity
18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Liu C,
Wang Z,
Zhou P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000US-0598042.
2000US-0620312.
2000US-0653450.
                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000US-0488725.
2000US-0552317.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000WO-US34263
                                     endocarditis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-0727344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000US-0662191.
2000US-0693036.
                                                                                    (first entry)
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                                                                genomic polynucleotide sequence SEQ ID NO:4307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Asundi V, Ch
Wehrman T, X
Goodrich R,
                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                    4.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         polypeptides, us
system injuries
                                      ds.
                                                                                                                                                                                                                0;
                                                                                                                                                                                                                          Score 18; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chen R,
Xu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nen R, Ma Y,
Ku C, Xue AJ,
Drmanac RT;
                                               strain;
                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   useful for treating disorders
                                                                                                                                                                                                                         DB
23;
                                               infection; diagnosis;
                                                                                                                                                                                                                                  22;
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Yang Y,
                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                  Length 3556;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zhang
                                                                                                                                                                                                                0,
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                                                                                                                                                                                                               Gaps
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RESULT 43
AAH54174/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                           Staphylococcus epidermidis vaccination; endocarditis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1763 TGCATCAGCTGCTGCAAC 1746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3633 BP; 1340 A; 498 C; 707 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 8; Page 2042-2043; 2188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acids encoding polypeptides from Staphylococcus epidermidis, useful for vaccinating against infections, e.g. endocarditis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
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09-NOV-1999;
                                                    09-NOV-2000; 2000WO-US30782
                                                                                                                                                                                                                       Staphylococcus
                                                                                                                                                                                                                                                                                                                                                               S. epidermidis
                                                                                                                                                                                                                                                                                                                                                                                                                    03-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH54174 standard; DNA; 4106
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                                                                                                           17-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        269 TGCATCAGCTGCTGCAAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                       epidermidis
99US-0164258
                                                                                                                                                                                                                                                                                                                                                               genomic polynucleotide sequence SEQ ID NO:3538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               286
                                                                                                                                                                                                                                                                                 SR1 strain; infection; diagnosis;
ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22;
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ABL13144/c
ID ABL13144 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH52304 to AAH53970 represent nucleic acids (I) encoding polypeptides (II), given in AAG81454 to AAG83120, from Staphylococcus epidermidis. (I) and (II) can have antibacterial activity and therefore can be used in vaccination. The nucleic acids (I) may be used to produce the CC sepidermidis polypeptides (II) via the production of vectors (I) containing them which are used to produce hosts cells which express the CC used to vaccinate subjects and to raise antibodies against the bacteria. The polypeptides (II) (and/or nucleic acids) may then be used to vaccinate subjects and to raise antibodies against the bacteria. The polypeptides may also be used to assay for other inhibitors of their creatment of S. epidermidis infections, e.g. endocarditis. AAH53971 to AAH55090 represent specifically claimed S. epidermidis genomic DNA CC polynucleotide sequences from the present invention. AAH55991 to CC AAH55098 represent oligonucleotide sequences and primers which are used in the exemplification of the present invention.

CC N.B. The present in the sequence listing of the present specification, thousand the sequence listing only goes up to SEQ ID NO:4454 so even thousand the present for SEQ ID NO:4455 to 4472, and so the present for SEQ ID NO:4465 to 4472, and so the present for SEQ ID NO:4465 to 4472, and so the present for SEQ ID NO:4465 to 4472, and so the present for SEQ ID NO:4465 to 4472, and so the present for SEQ ID NO:4465 to 4472, and so the present for SEQ ID NO:4465 to 4472, and so the present for SEQ ID NO:4465 to 4472, and so the present for SEQ ID NO:4465 to 4472, and so the present for SEQ ID NO:4465 to 4472.
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Best Local
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P-PSDB;
                  WPI; 2001-656860/75
                                                                                          (PEKE ) PE
                                                                                                                            23-MAR-2000;
11-JUL-2000;
                                                                                                                                                                               23-MAR-2001;
                                                                                                                                                                                                                   27-SEP-2001
                                                                                                                                                                                                                                                     WO200171042-A2
                                                                                                                                                                                                                                                                                         Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                         pharmaceutical; gene;
                                                                                                                                                                                                                                                                                                                                   Drosophila; developmental biology; cell signalling; insecticide
                                                                                                                                                                                                                                                                                                                                                                        Drosophila melanogaster expressed polynucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                      26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 4106 BP; 1436 A; 659 C; 643 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 no sequences are present for SEQ ID NO:4455 to 4464.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids encoding polypeptides from Staphylococcus epidermidis, useful for vaccinating against infections, e.g. endocarditis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    269 TGCATCAGCTGCTGCAAC 286
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                        CORP NY
                                                   Adams M,
                                                                                                                                                                               2001WO-US09231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                          2000US-191637P
2000US-0614150
                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cDNA; 6399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.9%; Dr.
100.0%; Pr.
0;
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                                                   , DWG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 18; DB; Pred. No. 24; 0; Mismatches
                                                   Myers EW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .
24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22;
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ABL15972/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local :
        capable of detecting 1000 or more genes from Drosophila. The invention useful in developmental biology and in elucidating cell signalling and cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention
discloses genomic DNA sequences (ABL16176-ABL30511),
                                                                 The invention relates to an isolated nucleic acid detection reagent
                                                                                                 Claim
                                                                                                                          New isolated nucleic acid genes from Drosophila and interactions -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5810
                                                                                                                                                                                                                           Venter
                                                                                                                                                                                                                                                    (PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                 23-MAR-2000;
11-JUL-2000;
                                                                                                                                                                                                                                                                                                                       23-MAR-2001; 2001WO-US09231
                                                                                                                                                                                                                                                                                                                                                                                WO200171042-A2
                                                                                                                                                                                                                                                                                                                                                                                                            Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                                                                                                               Drosophila; developmental biology; cell signalling; insecticide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Drosophila melanogaster expressed polynucleotide SEQ ID NO 42398.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL15972 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The invent useful in developmental biology and in elucidating cell signalling cell-cell interactions in higher enkaryotes for the development of
                                                                                                                                                                                                                                                                                                                                                      27-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                     pharmaceutical; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABL15972;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABL16176-ABL30511), expressed DNA sequences (ABL01840-ABL16175) and the encoded proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6399 BP; 1706 A; 1543 C; 1399 G; 1751 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID NO 33914; 21pp + Sequence Listing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                110 GTGAATTGCACGGTGAAC 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interactions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genes from Drosophila and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                 1;
                                                                                                                                                                                               2001-656860/75.
                                                                                                                                                                                                                           JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
                                                                                                                                                                                  ABB71869
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                                                                                                SEQ ID NO
                                                                                                                                                                                                                           Adams M,
                                                                                                                                                                                                                                                                                2000US-191637P
2000US-0614150
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                42398;
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                                                                                                                                                                                                                        PWD,
                                                                                            21pp + Sequence Listing; English
                                                                                                                                     detection reagent for detecting 1000 or more for elucidating cell signalling and cell-cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     detection reagent for detecting 1000 or more for elucidating cell signalling and cell-cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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Minimum DB
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seg length: 2000000000
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Match
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369
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Copyright (c) 1993 - 2002 Compugen
          /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/backfiles1.seq:*
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        15967
15967
13936
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  US-09-134-001C-1425
US-09-134-001C-2083
US-09-134-001C-263
US-09-221-0178-627
US-08-532-8288-12
US-08-532-8288-1
US-08-532-8288-7
US-08-532-8288-7
US-08-532-8288-9
US-08-532-8288-9
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US-08-700-359-8
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US-08-700-359-10
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US-08-700-359-10
US-08-967-104-7
US-08-985-908-3
US-08-985-908-3
US-08-985-908-3
US-08-852-730-13
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US-08-985-916-3
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8184.096 Million cell updates/sec
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Sequence 1425, Apsequence 263, App sequence 263, App sequence 17, Appli sequence 12, Appli sequence 1, Appli sequence 2, Appli sequence 2, Appli sequence 3, Appli sequence 10, Appli sequence 21, Appli sequence 5, Appli sequence 5, Appli sequence 21, Appli sequence 5, Appli sequence 5, Appli sequence 5, Appli sequence 6, Appli sequence 4, Appli sequence 12, Appli sequence 13, Appli sequence 3, Appli sequence 4, Appli sequence 3, Appli sequence 4, Appli sequence 3, Appli sequence 4, Appli sequence 
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RESULT 1  US-09-134 001  Sequence 14  Patent No.  GENERAL INF.  APPLICANT:  TITLE OF I  FILE REFER  CURRENT AP  CURRENT AP  CURRENT FILE  PRIOR APPLI  PRIOR APPLI  PRIOR FILL  TYPE: DNA		C C C C C C C C C C C C C C C C C C C
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		1643 1643 19307 19307 19307 19307 2932 1572 2502 2702 2702 37950 37950 37950 37950 37950 37950 37950 37950 37950
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		US-08-985-916-4 US-08-985-916-6 US-08-836-022A-10 US-09-427-048A-10 US-09-094-287-8 US-09-094-287-3 US-09-094-287-3 US-09-094-287-3 US-09-094-287-3 US-09-461-697-359 US-09-461-697-359 US-09-453-702B-258 US-09-338-907-183 US-09-218-207-183 US-09-318-207-183 US-09-318-207-183 US-09-318-207-183 US-09-453-702B-57 US-09-453-702B-57 US-09-453-702B-57 US-08-368-803-21 US-08-578-096A-22
		Sequence 4, Appli Sequence 6, Appli Sequence 10, Appl Sequence 10, Appl Sequence 11, Appli Sequence 1, Appli Sequence 1, Appli Sequence 2, Appli Sequence 2, Appli Sequence 258, App Sequence 183, App Sequence 183, App Sequence 17, Appli Sequence 17, Appli Sequence 27, Appli Sequence 27, Appli Sequence 27, Appli Sequence 27, Appli Sequence 21, Appli Sequence 21, Appli

Sequence 2083, Application US/09134001C
Patent NO. 6380370
GENERAL INFORMATION:
APPLICANT: Lynn DOUGETTE-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENTIAL SEQUENCE: GRO-007
FILE REFERENCE: GRO-007
CURRENT APPLICATION NUMBER: US/09/134,001C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 2083
LENGTH: 1596
TYPE: DNA ; TYPE: DNA ; ORGANISM: Staphylococcus epidermidis US-09-134-001C-1425 RESULT 2 US-09-134-001C-2083/c Ş Query Match Best Local Matches 567 Local Similarity 18; Conservative 4.9%; Score 18; 100.0%; Pred. No. 0; Mismatches AMINO ACID SEQUENCES RELATING DB 4; 0; Indels Length 1587; AND 0 Gaps TO STAPHYLOCOCC 0

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TYPE: DNA
; ORGANISM: Staphylococcus epidermidis
US-09-134-001C-263
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US-09-134-001C-263/C
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC-007
CURRENT APPLICATION NUMBER: US/09/134,001C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR TILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 263
LENGTH: 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: PP1182
FILING DATE: 31-DEC-1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                    ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: ROSS, BIUCE C.
TITLE OF INVENTION: P. GINGIVALIS NUCLEOTIDES AND USES THEREOF NUMBER OF SEQUENCES: 1120
CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER: PP1546 FILING DATE: 30-JAN-1998
                                                                                                          APPLICATION NUMBER: US/0 FILING DATE: 23-DEC-1998 CLASSIFICATION:
                                                                                                                                                                                                COMPUTER: IBM Compatible OPERATING SYSTEM: Windows
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              y Match 4.9%; So
Local Similarity 100.0%; P
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                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/09221017B
                                                                                                                                                                                                                                                                                                                                               755 PAGE MILL ROAD
                                                                                                                                                                                                                                                                                           USA
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Pred. No.
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Pred. No.
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US-08-532-828B-11/c
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Best Local Similarity
Matches 17; Conserv
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INFORMATION FOR SEQ ID NO:
              PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-101450
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: SUZUKI, Tomoko
APPLICANT: TANAKA, AKİKO
APPLICANT: MATSUI, HİROSİİ
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 27
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-813-5600
                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: SUGIMOTO, Masakazu APPLICANT: OGAWA, Yuri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORIGINAL SOURCE:
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HYPOTHETICAL:
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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                                                                                               APPLICATION NUMBER: US/08/532,828B FILING DATE: 27-OCT-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                   STREET: 1755 S. CITY: ARLINGTON
                                                                                                                                                                                                                                                                                          STATE:
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
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STRANDEDNESS: double
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FTIING DATE: 10-DEC-1998
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REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM:
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                                                                                                                                                                                                                                                                                                                     N, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. JEFFERSON DAVIS HIGHWAY, SUITE 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.6%; Score 17;
100.0%; Pred. No.
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24,618
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
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Best Local Similarity
Matches 17; Conserv
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: SUGIMOTO, Masakazu
APPLICANT: OGAWA, Yuri
APPLICANT: SUZUKI, Tomoko
APPLICANT: TANAKA, Akiko
APPLICANT: MATSUI, Hiroshi
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
NUMBER OF SEQUENCES: 24
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MOLECULE TYPE: (
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                       TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO: 12:
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LENGTH: 1263 nucleotides
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                                                                                                                                                                                         APPLICATION NUMBER: JP 5-10:
FILING DATE: 27-APR-1993
ATTORNEY AGENT INFORMATION:
NAME: NORWAN F. OBLON
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10
                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOTWARE: MS-DOS TEXT EDITOR CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 1263 nucleotides
                                                                                                                                                          TELEPHONE: 703-413-3000
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 HYPOTHETICAL:
                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: doub
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                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/532,828B FILING DATE: 27-OCT-1995 CLASSIFICATION: 435
                                                                                                                           TELEPHONE: 703-413-2220
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
                                  TOPOLOGY:
                                                 STRANDEDNESS: double
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                                                                nucleic acid
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NO
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; ORIGINAL SOURCE:
; ORGANISM: Corynebacterium glutamicum
; STRAIN: AJ3463
US-08-532-828B-12
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                                                                                                          HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Corynebacterium glutamicum
STRAIN: ATCC 13869
US-08-532-828B-1
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                                                 Matches
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Best Local S
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1643 nucleotides
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-101450
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS:
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: MATSUI, HİrOShİ
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
                                                                                                                                                                                                                                                                                                                                       NAME: NORMAN F. OBLON
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
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                                                                                                                                                                                                        MOLECULE TYPE:
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161 CAAAGTGCCGGGATCAT 177
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CITY: ARLINGTON
STATE: VA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/532,828B FILING DATE: 27-OCT-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                            TELEPHONE: /UJ 703-413-2220
                                                              Local Similarity
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                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                           TYPE: nucleic acid
                                                 17;
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OGAWA, Yuri
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                                                 Conservative
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                                                             4.6%; Score 17; DB
100.0%; Pred. No. 15;
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                                                                                 DB 1;
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936 CANAGTGCCGGGATCAT 920

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US-08-532-828B-7/c
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US-08-532-828B-2
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                                                                        Sequence 7, Application US/08532828B Patent No. 5688671
                                                                                                                                                                                                                               Matches
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                                                         GENERAL INFORMATION:
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                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-101450
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: OGAWA, Yuri
APPLICANT: SUZUKI, Tomoko
APPLICANT: TANAKA, Akiko
APPLICANT: MATSUI, Hiroshi
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
                APPLICANT:
                                                                                                                                                                                                                                                                                                                                  ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: ge
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                   161 CAAAGTGCCGGGATCAT 177
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                                                                                                                                                                  936 CAAAGTGCCGGGATCAT 920
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                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
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17; Conservative

    Application US/08532828B
    5688671

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        SUGIMOTO, Masakazu
OGAWA, Yuri
SUZUKI, Tomoko
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TANAKA, Akiko
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                                                                                                                                                                                                                                                            4.68;
                                                                                                                                                                                                                                          Score 17;
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                                                                                                                                                                                                                                            No.
                                                                                                                                                                                                                                          DB 1; Length 1643; 15;
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RESULT 10
US-08-532-828B-8/c
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                                                                                                                                                                                         APPLICANT: TANAKA, AKİKO
APPLICANT: MATSUI, HİROShİ
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
NUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: JP 5-101450 FILING DATE: 27-APR-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                             CORRESPONDENCE ADDRESS:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 1643 nucleotides
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
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MEDIUM TYPE: Floppy
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  936 CAAAGTGCCGGGATCAT 920
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LOCATION:
COUNTRY:
                                                           CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ' Match
Local Similarity 100.0%; P
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                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /U5 -- TELEPHONE: /
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                                                                                                                                 ADDRESSEE:
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                                                                                       E: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
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USA
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217..1482
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ZIP: 22202 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy

IBM PC compatible SYSTEM: PC-DOS/MS-DOS

Floppy disk

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US-08-532-828B-8
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US-08-532-828B-9/c
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APPLICATION NUMBER: JP 5-101450
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10-764-0
                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:

APPLICANT: SUGIMOTO, Masakazu
APPLICANT: OSAWA, Yuri
APPLICANT: SUZUKI, Tomoko
APPLICANT: TANAKA, Akiko
APPLICANT: TANAKA, Akiko
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TELEPHONE: 703-413-3000
TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                              APPLICANT: MATSUI, Hiroshi
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               161 CAAAGTGCCGGGATCAT 177
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/532,828B FILING DATE: 27-OCT-1995
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SOFTWARE: MS-DOS TEXT EDITOR
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CITY: ARLINGTON
                                                                                                                                                                COUNTRY: U
ZIP: 22202
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                       APPLICATION NUMBER:
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UMBER: US/08/532,828B
27-OCT-1995
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US-08-532-828B-9
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US-08-532-828B-10/c
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Best Local Similarity
Matches 17; Conserv
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INFORMATION FOR SEQ ID NO: 9
SEQUENCE CHARACTERISTICS:
LENGTH: 1643 nucleotides
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-1
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                  APPLICATION NUMBER: US/08/532,828B
FILING DATE: 27-OCT-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-101450
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
C. 18
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REFERENCE/DOCKET NUMBER: 10-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
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APPLICANT: OGAWA, YUIİ
APPLICANT: SUZUKI, Tomoko
APPLICANT: TANAKA, AKİKO
APPLICANT: TANAKA, AKİKO
APPLICANT: MATSUI, HİFOSİİ
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: BM PC COMPATION
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Corynebacterium glutamicum STRAIN: ATCC13869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: 703-413-2220
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                             COUNTRY: UZIP: 22202
                                                                                                                                                                                                                                                                                                                                                                  STATE:
                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
                REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid
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100.0%; Pred. No. 15;
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                    10-764-0 PCT
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TELEPHONE:

: 703-413-3000 703-413-2220

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US-08-532-828B-10
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                                                                               TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: SUGIMOTO, MASAKAZU
APPLICANT: USUDA, YOSHIHIRO
APPLICANT: SUZUKI, TOMOKO
APPLICANT: TANAKA, AKIKO
APPLICANT: MATSUI, HIROSHI
TITLE OF INVENTION: METHOD OF P
NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 10
SEQUENCE CHARACTERISTICS:
LENGTH: 1643 nucleotides
 MOLECULE TYPE:
                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 703-413-3000
                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP-3:
FILING DATE: 04-WAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
BEGISTERS NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-DOS/MS-I
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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HYPOTHETICAL:
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                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/700,359 FILING DATE: 08-OCT-1996
                   TOPOLOGY:
                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                     REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: ARLINGTON
                                                                 LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION:
                                               nucleic acid
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ative 0;
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100.0%; Pred. No.
                                                                                                                                                                               24,618
                                                                                                                                                                   10-819-0 PCT
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US-08-700-359-8
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US-08-700-359-8/c
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US-08-700-359-7
       Matches
                Best Local
                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 8, Application US/08700359 Patent No. 5766925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                            MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 04-MAR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: JP-35019
                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: MS-DOS TEXT EDITION
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/700
FILING DATE: 08-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORIGINAL SOURCE:
ORGANISM: Corynebacterium glutamicum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MATSUI, HIROSHI
TITLE OF INVENTION: METHOD OF PRODUCING L-LYSINE
NUMBER OF SEQUENCES: 22
                                                                                        NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              161 CAAAGTGCCGGGATCAT 177
                                                                         IDENTIFICATION METHOD:
                                                                                                                                 ORGANISM: Corynebacterium glutamicum STRAIN: ATCC13869
                                                                                                                                                                                                                                                            TELEPHONE: 703-413-2220
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                                                                                                                                                 ORGANISM:
                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 24
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 1755 S. CITY: ARLINGTON
                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE:
                                                                                                                                                                                                                                  CENGTH:
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17; Conserv
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    Conservative
                                                                                        mat peptide 217..1479
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                                                                                                                                                                                          linear
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                        703-413-3000
                                                                                                                                                                         genomic DNA
                                                                                                                                                                                                      double
                4.6%; Score 17; DB 1; 100.0%; Pred. No. 15;
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Pred. No.
    Mismatches
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                            Length 1643;
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   Indels
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0; Gaps
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161 CARAGTGCCGGGATCAT 177

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       US-08-674-168-21/c
; Sequence 21, Application US/08674168
; Patent No. 5804414
; GENERAL INFORMATION:
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                                                                                   RESULT 16
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GENERAL INFORMATION:
                                                                                                                                                                                                        Best Local Similarity Matches 17; Conserv
                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:

APPLICATION NUMBER: JP-35019

FILING DATE: 04-MAR-1994

ATTORNEY/AGENT INFORMATION:

NAME: NORMAN F. OBLON

REGISTRATION NUMBER: 24,618

REFERENCE/DOCKET NUMBER: 10-81:

TELECOMMUNICATION INFORMATION:

TELEPHONE: 703-413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: METHOD OF PRODUCING L-LYSINE
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C.
STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                936 CAAAGTGCCGGGATCAT 920
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APPLICANT: MORIYA, Mika
                                                                                                                                   936 CAAAGTGCCGGGATCAT 920
                                                                                                                                                                    161 CAAAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                                        NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/700,359 FILING DATE: 08-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                          STRAIN:
                                                                                                                                                                                                                                                                                                                                                                              ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                           LOCATION:
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ATCC13869
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USUDA, YOSHIHIRO
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                                                                                                                                                                                                    4.6%; Score 17; DB 1; llarity 100.0%; Pred. No. 15; Conservative 0; Mismatches
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964..1479
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                                                                                                                                                                                                                                                                                                                                                                                                                                               double
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; ORGANISM: Brevibact
; STRAIN: ATCC 13869
US-08-674-168-21
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US-08-596-366-5/c
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 413-3000
TELEFAX: (703) 413-2220
TELEEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                        Sequence 5, Application US/08596366 Patent No. 5876983
                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: METHOD OF AMPLIFYING GENE USING TITLE OF INVENTION: ARTIFICIAL TRANSPOSON NUMBER OF SEQUENCES: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 7-166541
FILING DATE: 30-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: OBLON, NORMAN F.
REGISTRATION NUMBER: 24,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ANTI-SENSE: NO ORIGINAL SOURCE:
               APPLICANT: IZUI, Katsura
TITLE OF INVENTION: MUTANT PHOSPHOENOLPYRUVATE CARBOXYLASE,
TITLE OF INVENTION: ITS GENE, AND PRODUCTION METHOD OF AMINO ACID
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                     APPLICANT: SUGINOTO, Masakazu
APPLICANT: SUZUKI, TOMOKO
APPLICANT: MATSUJI, Hiroshi
APPLICANT: IZUI, Katsura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                           161 CAAAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                        936 CAAAGTGCCGGGATCAT 920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 01 CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/674,168
                                                                                                                                                                                                                                                                                                                                                            Local Similarity
les 17; Conserv
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 ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1643 base pairs
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YOKOZEKI, Kenzo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IBM PC compatible
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OBLON, SPIVAK, McCLELLAND, MAIER, & NEUSTADT
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Pred. No.
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RESULT 18
US-08-596-366-7/c
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US-08-596-366-5
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                                                                                                                                                                                                                                                               Sequence 7, Application US/08596366 Patent No. 5876983
                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: SUGIMOTO, Masaka
APPLICANT: SUZUKI, Tomoko
APPLICANT: MATSUI, Hiroshi
APPLICANT: IZUI, Katsura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (703) 413-3000
TELEFAX: (703) 413-220
TELEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: JP 5-209776
FILING DATE: 24-AUG-1993
PRIOR APPLICATION DATA:
APPLICATION UMBER: JP 6-153876
FILING DATE: 05-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: OBLON, NORMAN F.
REGISTRATION NUMBER: 24.618
REFERENCE/DOCKET NUMBER: 10-784-0
TELECOMMUNICATION INFORMATION:
THE PENNER: 7010 A 11-2000
                                                                                     TITLE OF INVENTION: MUTANT PHOSPHOENOLPYRUVATE CARBOXYLASE,
TITLE OF INVENTION: ITS GENE, AND PRODUCTION METHOD OF AMINO ACID
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORIGINAL SOURCE:
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APPLICATION NUMBER: (
FILING DATE: 24-AUG-:
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MEDIUM TYPE: Floppy disk
ADDRESSEE: OBLOW, STATES HIGHWAY, SUITE # 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY:
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TOPOLOGY: li
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
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                                                                                                                                                               IZUI, Katsura
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                                                                                                                                                                                                                          SUGIMOTO, Masakazu
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217..1482
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                                                                  OBLON, SPIVAK, MCCLELLAND, MAIER, & NEUSTADT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA (genomic)
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N: 435
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RESULT 19
US-08-967-104-5/c
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TELEFAX: (703) 413-220
TELEX: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 1643 base pairs
                                                                                                                                                                                                                                                                                  Sequence 5, Application US/08967104 Patent No. 5919694
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity Matches 17; Conserv
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APPLICANT: SUGIMO
APPLICANT: SUZUKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: JI
FILING DATE: 24-AUG-1:
PRIOR APPLICATION DATA:
                                                                                                                             TITLE OF INVENTION: MUTANT PHOSPHOENOLPYRUVATE CARBOXYLASE,
TITLE OF INVENTION: ITS GENE, AND PRODUCTION METHOD OF AMINO ACID
NUMBER OF SEQUENCES: 12
                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN: ATCC13869 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10-784-0 PCT
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA (genomic) ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: JP 6-153876
FILING DATE: 05-UTL-1994
ATTORNEY/AGENT INFORMATION:
NAME: OBLON, NORMAN F.
                                                                                                                                                                                                                                                                                                                                                                                      936 CAAAGTGCCGGGATCAT 920
                                                                                                                                                                                                                                                                                                                                                                                                        161 CAAAGTGCCGGGATCAT 177
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APPLICATION NUMBER:
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MEDIUM TYPE: Floppy disk
                                 STREET: 1755 JE
CITY: ARLINGTON
STATE: VIRGINIA
COUNTRY: UZIP: 22202
                                                                                  ADDRESSEE:
                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
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                               VIRGINIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid
                                                            1755 JEFFERSON DAVIS HIGHWAY, SUITE # 400
                                                                                                                                                                                                        SUGIMOTO, Masakazu
SUZUKI, Tomoko
MATSUI, Hiroshi
                                                                                                                                                                                      IZUI, Katsura
                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.6%; Score 17; DB 2; ilarity 100.0%; Pred. No. 15; Conservative 0; Mismatches
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964..1482
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                                                                                               OBLON, SPIVAK, McCLELLAND, MAIER, & NEUSTADT
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SYSTEM: PC-DOS/MS-DOS
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COMPUTER READABLE FORM:

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; NAME/KEY:
; LOCATION:
US-08-967-104-5
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FILING DATE: 24-AUG-1993

PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 6-153876

FILING DATE: 05-JUL-1994

ATTORNEY/AGENT INFORMATION:
NAME: OBLOW, NORMAN F.
NAME: OBLOW, NORMAN F.
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10-784-0 PCT
TELECOMMUNICATION INFORMATION:
TELEFAX: (703) 413-3000
TELEX: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 5:
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                 Sequence 7, Application US/08967104
Patent No. 5919694
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 24-AUG-
PRIOR APPLICATION DATA:
                                                                                                                                       APPLICANT: SUGIMOTO, Masakazu
APPLICANT: SUZUKI, TOMOKO
APPLICANT: SUZUKI, TOMOKO
APPLICANT: MATSUI, HIROShi
APPLICANT: IZUI, Katsura
TITLE OF INVENTION: MUTANT PHOSPHOENOLPYRUVATE CARBOXYLASE,
TITLE OF INVENTION: ITS GENE, AND PRODUCTION METHOD OF AMINO
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
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ORIGINAL SOURCE:
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COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     161 CAAAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                                                                                                                                                     936 CAAAGTGCCGGGATCAT 920
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                                               STATE: VIRGINIA MCCLELLAND, MAIER, & NE STATE: VIRGINIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Corynebacterium glutamicum STRAIN: ATCC13869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                              COUNTRY:
                   22202
                                  USA
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SYSTEM: PC-DOS/MS-DOS
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217..1482
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                                                                                                                        OBLON, SPIVAK, McCLELLAND, MAIER, & NEUSTADT
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24-AUG-1993
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100.0%; Pred. No.
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TELEFAX: (703) 413-220
TELEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 1643 base pairs
                                                                                                                                                                                                                                                                                                                     Sequence 3, Application US/08985908
Patent No. 6004773
GENERAL INFORMATION:
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Best Local Similarity
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RAIOR APPLICATION DATA:
APPLICATION NUMBER: JP 6-1538

FILING DATE: 05-JUL-1994

ATTORNEY/AGENT INFORMATION:
NAME: OBLON, NORMAN F.
REGISTRATION UNMBER: 24,618

REFERENCE/DOCKET NUMBER: 10-7

TELECOMMUNICATION INFORMATION:
COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PatentIn Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,908
FILING DATE: 05-DEC-1997
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
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COUNTRY: UC.
-+D: 22202
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                                                          Release #1.0, Version #1.30
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CLASSIFICATION:

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TELECOMMUNICATION INFORMATION: 703-413-3000
                                                                                                                                                 TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO:
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                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 1643 bases
TYPE: nucleic acid
                            MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                               NAME: NORMAN F. OBLON
REGISTRATION NUMBER: 24,618
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
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APPLICATION NUMBER: JP 8-325659
FILING DATE: 05-DEC-1996
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                                                                                                                                                                                                                                                 FILING DATE: 05-DEC-1996 ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS
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LENGTH: 1643 bases
TYPE: nucleic acid
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               ORGANISM:
                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/985,908 FILING DATE: 05-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, FOURTH FLOOR CITY: ARLINGTON
                                                                 TOPOLOGY:
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ATCC 13869
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IVENTION: METHOD FOR PRODUCING L-LYSINE
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        Brevibacterium lactofermentum
                                                               linear
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SYSTEM: PC-DOS/MS-DOS
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TELEPAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
'ENGTH: 143 bases
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US-08-985-908-6/c
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Best Local Similarity
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ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
REGISTRATION UNMBER: 24,618
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
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PRIOR APPLICATION NUMBER: (
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936 CAAAGTGCCGGGATCAT 920
                                                              161 CAAAGTGCCGGGATCAT 177
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION: METHOD FOR PRODUCING L-LYSINE
NUMBER OF SEQUENCES: 31
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                                                                                                                                                                                                                                                                                                         LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN: ATCC 13869
                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 05-DE
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OPERATING SYSTEM: PC-DOS/MS-DOS
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les 17; Conserv
                                                                                                                                    17;
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                                                                                                                                    Conservative
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217..1482
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05-DEC-1996
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                                                                                                                                                                    100.0%;
                                                                                                                                                                4.6%; Score 17;
100.0%; Pred. No.
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                                                                                                                                                                                                DB 3;
                                                                                                                                                                                                Length 1643;
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US-08-852-730-12/c ; Sequence 12, Application US/08852730

RESULT 24

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RESULT 25
US-08-852-730-13/c
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Best Local Similarity
Matches 17; Conserva
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 12:
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APPLICANT: MASAKO IZUI, ATSUSHI HAYAKAWA, YASUHIKO YOSHIHARA, AND TSUYOSHI
APPLICANT: NAKAWATSU
TITLE OF INVENTION: METHOD OF PRODUCING L-LYSINE
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
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FILING DATE: 05-JUN-1996
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
REGISTRATION NUMBER: 24,618
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MEDIUM TYPE: Floppy disk
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LENGTH: 1643 base pairs
TYPE: nucleic acid
CTRANDEPONECS: 40mble
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ORIGINAL SOURCE:
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STATE: VA
22026
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSE: OBLON, SPIVAK, MCCLELLAND, MAIER AND NEUSTADT, P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY
CITY: ARLINGTON
                                                                                                                                                                                                                   TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                        APPLICANT: SEIKO HIRANO, MASAKAZU SUGIMOTO, EIICHI NAKANO, APPLICANT: MASAKO IZUI, ATSUSHI HAYAKAWA, YASUHIKO YOSHIHARA, AND TSUYOSHI APPLICANT: NAKAMATSU
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                                                                                   Cli.
STATE: va
STATE: 22026
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Brevibacterium lactofermentum STRAIN: ATCC 13869
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FILING DATE: 05-07-1997
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                                                                                                                                                                                                                                        METHOD OF PRODUCING L-LYSINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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                 TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER:
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ORIGINAL SOURCE:
ORGANISM: Brev
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LENGTH: 1643 base pair
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US/
FILING DATE: 05-07-1997
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                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                  APPLICATION NUMBER: JP 8-142812
FILING DATE: 05-JUN-1996
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
REGISTRATION NUMBER: 24,618
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SEQUENCE CHARACTERISTICS:
                                                                   TELECOMMUNICATION INFORMATION:
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TELEPHONE: 703-413-2220
TD NO:
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LOCATION:
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                 TELEPHONE: /v/
TELEPHONE: /v/
TO3-413-2220
TO TO NO:
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STATE: VA
ZIP: 22026
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CLASSIFICATION:
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1755 S. JEFFERSON DAVIS HIGHWAY
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                                                   703-413-3000
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                                                        Matches
                                                                                   Query Match
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICANT: ATSUSH
                                                                                                                          MOLECULE TYPE: q
ORIGINAL SOURCE:
ORGANISM: Brev
STRAIN: ATCC 1
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REGISTRATION INFORMATION:
TELEPHONE: 703-413-3000
                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: JP 8-325658
FILING DATE: 05-DEC-1996
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
936 CAAAGTGCCGGGATCAT 920
                          161 CARAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: (
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      161 CAAAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     936 CAAAGTGCCGGGATCAT 920
                                                                                                                                                                                                                                                       TELEPHONE: 703-413-2220
TO TO NO:
                                                                   Local
                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 05-DE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE:
                                                                                                                                                                                                                             LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
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                                                                   Similarity
                                                                                                                                                                                                            nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T: 1755 S. JEFFERSON DAVIS HIGHWAY, FOURTH FLOOR ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1: 1643 base pairs
                                                                                                                             ATCC 13869
                                                                                                                                                                                                                              1643 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          M: Brevibacterium lactofermentum
ATCC 13869
                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATSUSHI HAYAKAWA, MASAKAZU SUGIMOTO, YASUHIKO YOSHIHARA, AND TSUYOSHI
VENTION: METHOD FOR PRODUCING L-LYSINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                     PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                      Brevibacterium lactofermentum
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964..1482
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                    genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                             05-DEC-1997
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                                                                                                                                                                                                  double
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                                                                    100.0%;
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                          us/08/985,916
                                                     0
                                                                   Score 17; Pred. No.
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                                                  Mismatches
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                                                                 DB 4;
15;
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                                                    0; Indels
                                                                             Length 1643;
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US-08-985-916-4/c
                                                                                                                                                                         Sequence 6, Application US/08985916 Patent No. 6221636 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 4, Application US/08985916 Patent No. 6221636
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: ATSUSH
                                                                                               APPLICANT: ATSUSHI HAYAKAWA, MASAKAZU SUGIMOTO, YASUHIKO YOSHIHARA, AND TSUYOSH TITLE OF INVENTION: METHOD FOR PRODUCING L-LYSINE CORRESPONDENCES: 24
CORRESPONDENCE ADDRESS:
COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: genomic DNA
ORIGINAL SOURCE:
ORGANISM: Brevibacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 1643 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: JP 8-325658
FILING DATE: 05-DEC-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                        936 CAAAGTGCCGGGATCAT 920
                                                                                                                                                                                                                                                                                                                                          161 CAAAGTGCCGGGATCAT 177
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STREET: ARLINGTON
                                              STREET: 1755 S. CITY: ARLINGTON
                                 COUNTRY:
                                                            ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTAD' STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, FOURTH FLOOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 703-413-2220
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                              l Similarity
17; Conser
                    22152
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATSUSHI HAYAKAWA, MASAKAZU SUGIMOTO, YASUHIKO YOSHIHARA, AND TSUYOSH
VENTION: METHOD FOR PRODUCING L-LYSINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Brevibacterium lactofermentum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   703-413-3000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           double
                                                                                                                                                                                                                                                                                                                                                                                           4.6%; Score 17; DB 4; 100.0%; Pred. No. 15;
                                                                                SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Release #1.0, Version #1.30
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; LOCATION:
US-08-985-916-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10, Application US/08836022A Patent No. 6001557
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 6001557
GENERAL INFORMATION:
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APPLICATION NUMBER: JP 8-325658

FILING DATE: 05-DEC-1996

ATTORNEY/AGENT INFORMATION:

NAME: NORMAN F. OBLON

REGISTRATION NUMBER: 24,618

TELECOMMUNICATION INFORMATION:

TELEPHONE: 703-413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 1643 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Trustees of the University of Pennsylva APPLICANT: Wilson, James M.
APPLICANT: Fisher, Krishna J.
APPLICANT: Fisher, Krishna J.
APPLICANT: Weitzman, Matthew
APPLICANT: Weitzman, Matthew
TITLE OF INVENTION: Improved Adenovirus Virus and NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: genomic DNA ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                  FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/331,381
FILING DATE: 28-OCT-1994
                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Re-lease #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  161 CAAAGTGCCGGGATCAT 177
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE:
COMPUTER: IF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/985,916 FILING DATE: 05-DEC-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                         STREET: Spring House Corporate Cntr, CITY: Spring House
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN: ATCC 13869
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                       ZIP:
                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                               MEDIUM TYPE:
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                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                     19477
                                                                                                                                                                                                                                                                       USA
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964..1482
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trustees of the University of Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Brevibacterium lactofermentum
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                                                                                                                                                                                                           Floppy disk
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100.0%; Pred. No.
                                                                                                                    US/08/836,022A
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. 15;
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US-08-836-022A-10
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US-09-427-048A-10/c
US-09-427-048A-10
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 19307 base pai
                                                                                                                             REFERENCE/DOCKET NUMBER: GNVPN.008PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9200
TELEPAX: 215-540-5818
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9200
TELEFAX: 215-540-5818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3516 AAGAAGTGATGGAGCAA 3500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Bak, Mary E. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 19307 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOGTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 19477
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Improved Adenovirus Virus and
                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/836,022
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Howson and Howson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                 TOPOLOGY: unknown MOLECULE TYPE: cDNA
                 SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/427,048A FILING DATE: 21-Oct-1999 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Spring House
STATE: Pennsylvania
                                                                       LENGTH: 19307 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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Fisher, Krishna J.
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100.0%; Pr
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100.0%; Pred. No.
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                                                                                                                                  Sequence 1, Application US/09094287 Patent No. 6084071
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                                                                                   GENERAL INFORMATION:
APPLICANT: Racie, Lisa
APPLICANT: Jacobs, Kenr
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                 APPLICANT: Jacobs, Kenneth
TITLE OF INVENTION: Human L105 Proteins and Polynucleotide
TITLE OF INVENTION: Encoding Same
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: 11
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LENGTH: 535 base pairs
TYPE: nucleic acid
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TITLE OF INVENTION: Human L105 Proteins and Polynucleotide
TITLE OF INVENTION: Encoding Same
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      ADDRESSEE:
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les 16; Conserv
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REGISTRATION NUMBER: 32,724
REFERENCE/DOCKET NUMBER: GI
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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87 CambridgePark Drive
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617-876-5851
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Genetics Institute, Inc
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US-09-094-287-3/c
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                                      CUMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V6
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/094,287
FILING DATE:
CLASCITYTICATION OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY OF PROPERTY 
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APPLICANT: Jacobs, Kenneth
APPLICANT: Jacobs, Kenneth
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TELEFAX: 617-876-5851
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
ATTORNEY/AGENT INFORMATION
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ADDRESSEE: Genetics Institute,
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-498-8224
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SOSTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 87 Cambridge
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LOCATION:
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STRANDEDNESS: double
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Local Similarity 100.0%; P
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80..481
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human L105 Proteins and Polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Encoding Same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.3%; Score 16;
100.0%; Pred. No.
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                                                                                                                                                                 Version #1.30
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; LOCATION: (1)..(1569)
US-09-612-964-1
                                                                     RESULT 36
US-08-062-472B-2
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SEQUENCE CHARACTERISTICS:
LENGTH: 932 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
               Sequence 2, Application US/08062472B Patent No. 5695954 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 12
SEQ ID NO 1
LENCTH: 1572
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Best Local Similarity
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APPLICANT:
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APPLICANT:
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CURRENT FILING DATE: 2000-07-10
PRIOR APPLICATION NUMBER: RU 99114325
PRIOR FILING DATE: 1999-07-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Lunts Maria Grigorievna
APPLICANT: Kozlov Yury Ivanovich
APPLICANT: Ivanovskaya Lirina Valerievna
APPLICANT: Voroshilova Elvira Borlsovna
TITLE OF INVENTION: DNA CODING FOR MUTANT ISOPROPYLMALATE SYNTHASE,
TITLE OF INVENTION: L-LEUCINE-PRODUCING MICROORGANISM AND METHOD FOR PRODUCING
TITLE OF INVENTION: L-LEUCINE-PRODUCING MICROORGANISM AND METHOD FOR PRODUCING
TITLE REFERENCE: 193845USO
                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Escherichia coli
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TELEPHONE: 617-498-8224
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APPLICANT:
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                                                                                                                                                 547 ATCATCAGCGGCCTGT 562
                                                                                                                                                                                  196 ATCATCAGCGGCCTGT 211
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Brown, Scott A. REGISTRATION NUMBER:
                                                                                                                                                                                                                       watch 4.3%; Score 16; DB Local Similarity 100.0%; Pred. No. 49 nes 16; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gusyatiner Mikhail Markovich
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85..519
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154..519
Sherwood, Nancy G M
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; Pred. No. 49;
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. 49;
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: COGENT NEUROSCIENCE, Inc.
APPLICANT: LO, Donald C.
APPLICANT: Barney, Shawn
APPLICANT: Barney, Shawn
APPLICANT: Thomas, Mary Beth
APPLICANT: Portbury, Stuart D.
APPLICANT: Portbury, Stuart D.
APPLICANT: Ratz, Lawrence C.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING
TITLE OF INVENTION: AND TREATING CONDITIONS, DISORDERS, OR DISEASES INVOLVING
TITLE OF INVENTION: CELL DEATH
FILLE REFERENCE: 10001-005-999
CURRENT APPLICATION NUMBER: US/09/461,697
CURRENT FILING DATE: 1999-12-14
NUMBER OF SEQ ID NOS: 466
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 359
LEENGTH: 2702
TYPE: DNA
ORGANISM: Homo Sapiens
US-09-461-697-359
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US-09-461-697-359
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TELEPHONE: (503) 226-7391
TELEPAX: (503) 228-9446
INFORMATION FOR SEQ ID NO: 2:
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APPLICANT: McRory, John E
APPLICANT: Lescheid, David W
TITLE OF INVENTION: DNA ENCODING TWO FISH NEUROPEPTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: POLLEY, RICHARD J
REGISTRATION NUMBER: 2810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 14-MAY-19
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LENGTH: 2502 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                268 TIGCATCAGCIGCTGC 283
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 97204-2988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: KLARQUIST, SPARKMAN, CAMPBELL, LEIGH & ADDRESSEE: WHINSTON, LLP
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100.0%;
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                                                                                   RESULT 39
US-09-338-907-183/c
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US-09-453-702B-258
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                               Sequence 183, Application US/09338907 Patent No. 6265546 GENERAL INFORMATION:
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GENERAL INFORMATION:
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APPLICANT: Cohen, Daniel APPLICANT: Blumenfeld, Marta
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                                                                                                                                                          371 CCCAAGAAAAGGGGAA 386
                                                                                                                                                                                           311 CCCAAGAAAAGGGGAA 326
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Seay, Nicholas J.
NAME: Seay, Nicholas J.
REGISTRATION NUMBER: 27386
REFERENCE/DOCKET NUMBER: 960
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 251-5000
TELEFAX: (608) 251-9166
                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Diskette, 3.50 inch. 1.44Mb storage COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Word Perfect 8.0 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/453,702B
FILING DATE: 03-Dec-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/110,955
FILING DATE: 04-DEC-1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: <Unknown> PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                              16;
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                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1 South Pinckney Street CITY: Madison
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ZIP: 53701-2113
                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 8729 base pairs
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                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                             100.0%;
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                4.38;
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Pred. No.
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50;
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RESULT 40
US-09-218-207-183/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-338-907-183
                                                                                         GENERAL INFORMATION:
APPLICANT: Cohen, Daniel
APPLICANT: Blumenfeld, Marta
APPLICANT: Ilya, Chumakov
APPLICANT: Ilya, Chumakov
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: Prostate cancer gene
FILE REFERENCE: GENSET.018CP1
                                                                                                                                                                                                                              Sequence 183, Application US/09218207 Patent No. 6346381
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 183
LENGTH: 37950
                CURRENT APPLICATION NUMBER: US/09/218,207
CURRENT FILING DATE: 1998-12-22
EARLIER APPLICATION NUMBER: 08/996,306
EARLIER FILING DATE: 1997-12-22
EARLIER FILING DATE: 1997-1: EARLIER APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EARLIER APPLICATION NUMBER: 08/996,306
EARLIER FILING DATE: 1997-12-2
EARLIER APPLICATION NUMBER: 60/099,658
EARLIER FILING DATE: 1998-09-09
EARLIER APPLICATION NUMBER: 09/218,207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/338,907 CURRENT FILING DATE: 1999-06-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Ilya, Chumakov
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: PROSTATE CANCER GENE
FILE REFERENCE: GENSET.18CP1CP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patent.pm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EARLIER FILING DATE: 1998-12-22 NUMBER OF SEQ ID NOS: 578
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: exon
LOCATION: 25520..26016
OTHER INFORMATION: exon8
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NAME/KEY: exon
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LOCATION: 1462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: exon
LOCATION: 23387..23510
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LOCATION: 21789..21950
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LOCATION: 1267
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LOCATION: 1982
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                                                                                                                                                                                                                                                                                                                                                                                      142 TCAGAAAGAAGTGATG 157
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100.0%; Pred. No.
60/099,658
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EARLIER FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 578
SOFTWARE: Patent.pm
SEQ ID NO 183
LENGTH: 37950

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; SOFTWARE: PatentIn Ver. 2
; SEQ ID NO 17
; LENGTH: 48974
; TYPE: DNA
; ORGANISM: Mus musculus
US-08-920-422-17
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Db 43070 TGTCAGAAAGAAGTGA 43085
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: LOCATION: 25520..26016
: OTHER INFORMATION: exon8
US-09-218-207-183
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Db 34677 TCAGAAAGAAGTGATG 34662
                                                                                                                                                                                                                                                                                                                                             Sequence 17, Application US/08920422A Patent No. 6255473
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.0%;
watches 16; Conservative
                                                                                              Query Match
                                                                Matches
                                                                                                                                                                                                                 APPLICANT: Vitek, Michael P.
APPLICANT: Mitsuda, No. 6255473iaki
APPLICANT: ROSES, Allen D.
TITLE OF INVENTION: Presenilin-1 Gene Promoter
FILE REFERENCE: VITEKPRESENTLIN
CURRENT APPLICATION NUMBER: US/08/920,422A
CURRENT FILING DATE: 1997-08-29
NUMBER OF SEQ ID NOS: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: exon
LOCATION: 5259..5328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: exon7
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LOCATION: 23387..23510
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LOCATION: 21789..21950
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LOCATION: 14621..14710
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                             140 TGTCAGAAAGAAGTGA 155
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LOCATION: 12675..12791
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               142 TCAGAAAGAAGTGATG 157
                                                      Local Local 16;
                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19822..19912
                                                               Conservative
                                                          4.3%; Score 16;
100.0%; Pred. No.
tive 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.3%; Score 16; DB 4; 00.0%; Pred. No. 50;
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                                                           Mismatches
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Qy 45 AAATCCAGTGCTACCA 60
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Db 42450 AAATCCAGTGCTACCA 42465
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Patent No. 6087485
GENERAL INFORMATION:
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                                                                                                                                                                                                    Sequence 57, Application US/09453702B Patent No. 6365723 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/009,913
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 72928 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 36,677
REFERENCE/DOCKET NUMBER: SE
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Asthma Related Ger
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
ADDRESSEE: BOZICEVIC & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION: NAME: Sherwood, Pamela J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: AxyS Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Sherwood, Pamela REGISTRATION NUMBER: 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Quarles & Brady
STREET: 1 South Pinckney Street
                                                                                  TITLE OF INVENTION: No. 6365723el Sequences of E. coli 0157 NUMBER OF SEQUENCES: 265
                                                                                                                                                                                  APPLICANT: Blattner, Frederick R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid
CITY: Madison
STATE: WI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CA
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                                                                                                                                   Burland,
Burland,
Nicole T.
                                                                                                            Perna,
Plunkett, bu,
Ah. Rod
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                                                                                                                                                                                                                                                                                                                                                                                                            4.3%; Score 16; DB 3; 100.0%; Pred. No. 50;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch. 1.44Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Word Perfect 8.0

ZIP: 53701-2113

Fri Nov

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TOPOLOGY: Inear
; MOLECULE TYPE: DNA (genomic)
; SEQUENCE DESCRIPTION: SEQ ID NO: 57:
US-09-453-702B-57
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US-08-368-803-21
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GENERAL INFORMATION:
APPLICATION
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TELEFAX: (608) 251-9166
INFORMATION FOR SEQ ID NO: 57:
                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: LAPLACE, Eliane L
APPLICANT: RIVIERE, Michel A
APPLICANT: RIVIERE, Michel A
TITLE OF INVENTION: Avian Herpesvirus-based live recombinant avian
TITLE OF INVENTION: vaccine, in particular against Gumboro disease
NUMBER OF SEQUENCES: 27
                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      311 CCCAAGAAAAGGGGAA 326
                                   ATTORNEY/AGENT INFORMATION: NAME: SARRO, Thomas
                 TELECOMMUNICATION INFORMATION:
                                                                     APPLICATION NUMBER: US/01 FILING DATE: 05-JAN-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                              STREET: 727 SOUTH 23RD STREET CITY: ARLINGTON
                                                                                                                                                                                                                                                                 COUNTRY:
TELEPHONE:
                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/110,955
FILING DATE: 04-DEC-1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/453,702B
FILING DATE: 03-Dec-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                 22202
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 87563
                                                                                                                                                                                                                                                                                   VIRGINIA
                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DARTEIL, Raphael J
DUINAT, Carole V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BUBLOT, Michel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AUDONNET, Jean-Christophe F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                        LARSON AND TAYLOR
(703) 920-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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                                                                                                            US/08/368,803
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BER: 960296.95017
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US-08-578-096A-22
Search completed: November 7, 2002, 23:12:13 Job time: 123.827 secs
                                                       Вþ
                                                                                                                                                                         US-08-578-096A-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22, Application US/08578096A Patent No. 5980906 GENERAL INFORMATION:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (703) 892-8428
INFORMATION FOR SEQ ID NO: '.'
SEQUENCE CHARACTECS:
LENGTH: 36 base pairs
TYPE: nucleic acid
GTBANNERNEES: 5:5-1
                                                                                                                                            Query Match
                                                                                                                 Matches
                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              194 GCATCATCAGCGGCC 208
                                                                                    194 GCATCATCAGCGGCC 208
                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
les 15; Conserv
                                                                                                                                                                                                                               TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/578,096A
                                                                                                                                                                                                                  STRANDEDNESS: single
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                                                                                                                              Local
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100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Avian herpesvirus-based live recombinant
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Pred. No.
                                                                                                                                Score 15; pred. No.
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                                                                                                                    Mismatches
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hes 0;
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Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Post-processing: Listing first 45 summaries
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Maximum DB seq length: 200000000
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pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Published_Applications_NA: *
                                                                                                                                    .: //cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
.: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
.: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
.: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
.: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
.: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
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Copyright (c) 1993 - 2002 Compus
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8163.686 Million cell updates/sec
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## SUMMARIES

18 1	17 1	16 1	15 1	14 1	13 17	c 12 1	11 1	10 1	9 1	8	7 11	6 27	5 28	4 31	3 31	2 31	1 36	Result No. Score
6 4.6	.7 4.6	.7 4.6	.7 4.6	7 4.6	.7 4.6	.8 4.9	8 4.9	8 4.9	8 4.9	5 14.9	7 31.7			5 85.4			9 100.0	Query e Match
11842	8033	6252	2649	1713	1298	1791	286	256	237	430	480	1010	396	1897	1619	1619	369	Length DB
10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	BB
US-09-764-860-1187 US-09-815-242-2587	US-09-070-927A-121	US-09-964-824A-313	US-09-815-242-6571	US-09-815-242-7853	US-09-925-300-682	US-09-416-384A-6	US-09-294-093B-5021	US-09-294-093B-5709	US-09-294-093B-5287	US-09-867-701-4251	US-09-825-294-213	US-09-825-294-212	US-09-825-294-9	US-09-825-294-214	US-09-825-294-211	US-09-825-294-205	US-09-825-294-199	ID
Sequence 1187, Ap Sequence 2587, Ap	Sequence 121, App	Sequence 313, App	Sequence 6571, Ap	Sequence 7853, Ap	Sequence 682, App		Sequence 5021, Ap		Sequence 5287, Ap		Sequence 213, App	•	Sequence 9, Appli	Sequence 214, App	Sequence 211, App	Sequence 205, App	Sequence 199, App	Description

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3294 11749 17581	3212	2165	1989 2053	1549	1179	1172	689	654	615	538	488	414	411	406	405	402	375	369	364	271	236
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US-09-764-869-2170	US-09-922-261-359 US-09-834-291-1	US-09-802-669-94	US-09-801-368-41 US-09-822-849A-343	US-09-813-358-34	US-09-815-242-8625	US-09-815-242-4715	US-10-001-887-14	us-09-833-381-94	US-09-764-864-423	US-09-393-634-22	US-09-864-761-10902	US-09-983-965-3483	US-09-960-352-11005	US-09-960-352-3690	US-10-103-859-1	US-10-057-275-3	US-09-770-791-278	US-09-960-352-11486	us-09-960-352-1111	US-09-878-574-7775	US-09-923-876-3098
Sequence 2431, Ap Sequence 2170, Ap			Sequence 41, App1 Sequence 343, App	Sequence 34, Appl		Sequence 4715, Ap	Sequence 14, Appl	Sequence 94, Appl	Sequence 423, App	Sequence 22, Appl	Sequence 10902, A	Sequence 3483, Ap	Sequence 11005, A	Sequence 3690, Ap	Sequence 1, Appli	Sequence 3, Appli	Sequence 278, App	Sequence 11486, A	•	•	Sequence 3098, Ap

## ALIGNMENTS

```
APPLICANT: Study, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE REFERENCE: 210121.484C5
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 199
LENGTH: 369
                                                                                                                                                                                                                                                                  ; NAME/KEY: misc_feature
; LCCATION: (1)...(369)
; OTHER INFORMATION: n = A,T,C or G
US-09-825-294-199
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US-09-825-294-199
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                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0%;
                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
121 GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 180
                                                             369;
                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                   0;
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Pred. No. 4.5e-188;
D; Mismatches 0;
                                                                                                                                                                                                                                    Length 369;
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0,

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APPLICANT: X. Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Algate, Paul A.

APPLICANT: Algate, Paul A.

APPLICANT: Algate, Paul A.

APPLICANT: Filng, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA

FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 205

LENGTH: 1619
US-09-825-294-211

; Sequence 211, Application US/09825294

; Patent No. US20020004491A1

; GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
US-09-825-294-205
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                            GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 269
                                                                                                                           CTGCCTCGGCCCTCA 344
                                                                                                                                                                                                            GCATCAGCTGCTGCAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAAGGGGAAGTT 329
                                                                                                                                                              CTGCCTCGGCCCTCA 344
                                                                                                                                                                                           GCATCAGCTGCTGCAACACCCCCTCTTTGTAACGGGCCCAAGGCCCAAGAAAAGGGGGAAGTT 329
                                                                                                                                                                                                                                                                                                                                   AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            85.4%; Score 315; DB 10; 100.0%; Pred. No. 3.7e-159; vative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 1619;
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: XU, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 214
LENGTH: 1897
TYPE: DNA
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; Patent No. US20020004491A1
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-825-294-211
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APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Filing, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 211
FENCTH. 1610
   Query Match
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Best Local
                                                                                                                                                  NAME/KEY: misc_feature LOCATION: (1)...(1897)
                                                                                                                                                                                                                             ORGANISM: Homo sapiens FEATURE:
                                                                                                                  OTHER INFORMATION: n -
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                                                                                                              A,T,C or
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Patent No. US20020004491A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Filng, Steven P.
APPLICANT: Filng, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 215
SOFTWARE: FRSTSEQ for Windows Version 3.0
SEQ ID NO 9
LENGTH: 396
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Best Local Similarity
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LOCATION: (1)...(396)
OTHER INFORMATION: n = A,T,C or G
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                                                                                      AAGTGATGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
                                                                                                                                                    GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
                                                                   AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                   GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
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                                                                                                    GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION UMBER: US/09/825,294
CURRENT ETLING DATE: 2001-04-03
                               NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 213
LENGTH: 480
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        ORGANISM: Homo
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APPLICANT: Algate, Paul A.

APPLICANT: Filing, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN C.

FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 212

LENGTH: 1010
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                         271 CATCAGCTGCTGCAACACCCCTCTTTGTAAC 301
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US-09-294-093B-5287

; Sequence 5287, Application US/09294093B

; Patent No. US20010051335A1
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                                                                                                                              SOFTWARE:
SEQ ID NO 5287
                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/294,093B CURRENT FILING DATE: 1999-04-16 PRIOR APPLICATION NUMBER: 60/082,567 PRIOR FILING DATE: April 21, 1998 NUMBER OF SEQ ID NOS: 6207
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                                                                                                                                                                                                                                                         APPLICANT: Sherman, Bradley, K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN TASSEL
FILE REFERENCE: PL-0009 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Aglate, Paul A. APPLICANT: Jones, Robert
                                                                                                                                                                                                                                                                                                                          APPLICANT: Ito, Laura,
                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. US20010051335A1 700356180H1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapien
                                                       FEATURE:
                                                                     ORGANISM: Zea mays
                                                                                            TYPE: DNA
                                                                                                              ENGTH: 237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCAT 197
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                                                                                                                                                 PERL Program
                                                                                                                                                                                                                                                                                                                                           Lalgudi, Raghunath, V.
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100.0%; Pred. No. 5.6e-20;
tive 0; Mismatches 0;
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US-09-294-093B-5709
                                                                                      SOFTWARE:
SEQ ID NO 5021
LENGTH: 286
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                                                                                                                                                                                                                                                                                                                          Sequence 5021, Application Patent No. US20010051335A1 GENERAL INFORMATION:
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                                                                                                                                                                                            FILE REFERENCE: PL-0009 US
CURRENT APPLICATION NUMBER: US/09/294,093B
CURRENT FILING DATE: 1999-04-16
                                                                                                                                                                                                                                             APPLICANT: Lalgudi, Raghunath, V.
APPLICANT: Ito, Laura, Y.
APPLICANT: Sherman, Bradley, K.
APPLICANT: Sherman, Bradley, K.
TITLE OF INVENTION: POLYMUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN TASSEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                           PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: April
NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/294,093B CURRENT FILING DATE: 1999-04-16
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TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN TASSEL
FILE REFERENCE: PL-0009 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: PRIOR FILING DATE: April
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                                                    ORGANISM: Zea mays
                                                                           TYPE: DNA
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LOCATION: 19, 21, 234, 245
OTHER INFORMATION: a, t, c, g, or other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. US20010051335A1 700355590H1
                                       FEATURE:
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TYPE: DNA
ORGANISM: Zea mays
                  NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                Application US/09294093B
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                                                                                                                            PERL Program
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                                                                                                                                                            April 21,
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100.0%; Pred. No.
                                                                                                                                                                               60/082,567
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Pred. No.
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CURRENT APPLICATION NUMBER: US/09/925,300
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05988
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1890
SOFTWARE: PATENTIN VET: 2.0
SEQ ID NO 682
LENGTH: 1298
TYPE: DNA
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SEQ ID NO 6
LENGTH: 1791
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Best Local
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APPLICANT:
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CURRENT FILING DATE: 1999-10-12
CURRENT APPLICATION NUMBER: 05/09/416,384A
PRIOR APPLICATION NUMBER: 60/106,457
PRIOR FILING DATE: 1999-10-30
PRIOR APPLICATION NUMBER: 60/103,955
PRIOR APPLICATION NUMBER: 60/103,955
PRIOR FILING DATE: 1998-10-12
PRIOR FILING DATE: 1998-10-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1999-05-03
NUMBER OF SEQ ID NOS: 71
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TITLE OF INVENTION: Genes, proteins and biallelic markers related to central...
FILE REFERENCE: GENEET.045A0.
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LOCATION: 8, 26, 57, 117, 124, 216, 248,
OTHER INFORMATION: a, t, c, g, or other
                                                                                                                                                                                                                                                      APPLICANT: Craig Rosen,
APPLICANT: Steve Ruben
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA101
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ORGANISM: Homo sapiens
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nes 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                84 ACGACTGCTCCTCCCCG 101
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; NAME/RPY: misc feature
LOCATION: (1294)
; OTHER INFORMATION: n equals a,t,g, or c
US-09-925-300-682
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FILE REFERENCE: ELITRA.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
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PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-23
PRIOR FILING DATE: 2000-10-23
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                                                                                                                                                                                                                                                                                                                                                                                  ; NAME/KEY: CDS
; LOCATION: (1)...(1713)
US-09-815-242-7853
                                                                                                               RESULT 15
US-09-815-242-6571
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APPLICANT: Haselbeck, Robert
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 17; Conserv
Sequence 6571, Application US/09815242
Patent No. US20020061569A1
GENERAL INFORMATION:
APPLICANT: Haselbeck, Robert
APPLICANT: Ohlsen, Karl L.
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SEQ ID NO 7853
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PRIOR FILING DATE: 2000-11-7
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
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NUMBER OF SEQ ID NOS: 14110
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 1713
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Xu, H. Howard
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100.0%; Pred. No. 13;
tive 0; Mismatches 0
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Pred. No.
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. 13;
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Zyskind, Judith W.

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; TYPE: DNA; Homo sapiens US-09-964-824A-313
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; LOCATION: (1)...(2649)
US-09-815-242-6571
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APPLICANT: HORTIGAN, Stephen
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using SignaturiTLE OF INVENTION: Sets
FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT FILING DATE: 2001-09-27
CURRENT FILING DATE: 7001-09-27
Query Match 4.6%; Score 17; DB 10; Length 6252; Best Local Similarity 100.0%; Pred. No. 13; Matches 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                    SEQ ID NO 313
LENGTH: 6252
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PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
UNMBER OF SEQ ID NOS: 583
                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.0
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PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
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PRIOR FILING DATE: 2001-02-16
NUMBER OF SEQ ID NOS: 14110
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TITLE OF INVENTION: Identification of Essential Genes
TITLE OF INVENTION: Prokaryotes
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                                                                                                                                                                  LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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DR FILING DATE: 2000-10-23
DR APPLICATION NUMBER: 60/253,625
DR FILING DATE: 2000-11-27
DR APPLICATION NUMBER: 60/257,931
DR FILING DATE: 2000-12-22
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. 13;
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RESULT 18
US-09-764-860-1187
; Sequence 1187, Application US/09764860
; Patent No. US20020094953A1
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US-09-070-927A-121
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                                                                                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                 Matches
               APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC008
CURRENT APPLICATION NUMBER: US/09/764,860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 121
SEQUENCE CHARACTERISTICS:
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nes 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Kenley K. HOOVET REGISTRATION NUMBER: 40,302 REFERENCE/DOCKET NUMBER: PB TELECOMMUNICATION INFORMATION: TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: ASCII Text CURRENT APPLICATION DATA:
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Patrick J. Dillon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/046,655
FILING DATE: 1997-05-16
APPLICATION NUMBER: 60/044,031
FILING DATE: 1997-05-06
APPLICATION NUMBER: 60/066,009
FILING DATE: 1997-11-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
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TELEFAX: (301) 30
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OPERATING SYSTEM: MSDOS version 6.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Human Genome Sciences, Inc
STREET: 9410 Key West Avenue
                                                                                                                                                                                                                                                                                                                              Conservative
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APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
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; Sequence 2587, Application US/09815242
; Patent No. US20020061569A1
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                RESULT 20
US-09-923-876-3098/c
Sequence 3098, Application US/09923876
; Patent No. US20020013958A1
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                                                                                                                                                                                                                                                                                                     ; TYPE: DNA ; ORGANISM: Staphylococcus aureus US-09-815-242-2587
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SEQ ID NO 1187
LENGTH: 11842
TYPE: DNA
ORGANISM: Homo sapiens
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Best Local Similarity
Matches 17; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2587
LENGTH: 124
GENERAL INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/269,308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Haselbeck, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6500 CCTTCTGCTCCCCAGGG 6516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    237 CCTTCTGCTCCCCAGGG 253
                                                                                                                                          41 AATCCAGTGCTACCAG 56
                                                                                                                                                                                46 AATCCAGTGCTACCAG 61
                                                                                                                                                                                                                        Local Similarity
nes 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wall, Daniel
Trawick, John D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Carr, Grant J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ohlsen, Kari
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zyskind, Judith W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yamamoto, Robert T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                           4.3%; Score 16;
100.0%; Pred. No.
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100.0%; Pred. No.
tive 0; Mismatc
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                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                 DB 10;
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                                                                                                                                                                                                                                                              Length 124;
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                                                                                                                                                                                                                          Indels
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US-09-923-876-3098
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; OTHER INFORMATION: Clone ID: 701100324H1
US-09-878-574-7775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-878-574-7775
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                                                                                                                                                                                US-09-960-352-1111
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                                                                                                                                                                                                  RESULT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: La ROSA, Thomas J.

APPLICANT: Thompson, Michael D.

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with

TITLE OF INVENTION: Plants

FILE REFERENCE: 38-21(15401)B

CURRENT APPLICATION NUMBER: US/09/878,574

CURRENT FILING DATE: 2001-12-21

PRIOR APPLICATION NUMBER: 09/333,535

PRIOR APPLICATION NUMBER: 09/333,535

PRIOR FILING DATE: 1999-06-14

NUMBER OF SEQ ID NOS: 15775

SEQ ID NO 7775

LENGTH: 271
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Matches
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SEQ ID NO 3098
LENGTH: 236
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 7775, Application US/09878574 Patent No. US20020110548A1
GENERAL INFORMATION:
APPLICANT: Wairen, Wesley C.
APPLICANT: Tao, Nengbing
APPLICANT: Byatt, John C.
APPLICANT: Mathialagan, Nagappan
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
                                                                                                                                 sequence 1111, Application US/09960352
Patent No. US20020137139A1
                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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PRIOR FILING DATE: 1998-05-05
NUMBER OF SEQ ID NOS: 6332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/923,876
CURRENT FILING DATE: 2001-08-06
PRIOR APPLICATION NUMBER: 09/298,329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Kamigaki, Laura Y. (Ito)
APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN SEEDLING
FILE REFERENCE: PL-0012-1 CON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1999-04-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Byrum, Joseph R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. US20020013958A1 700161660H1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             273 TCAGCTGCTGCAACAC 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   131 TCAGCTGCTGCAACAC 116
                                                                                                                                                                                                                                                                                                             228 GGTACCAGTCCTTCTG 243
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                                                                                                                                                                                                                                                                                                                                                         16;
                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                           0;
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OTHER INFORMATION: Clone ID: 49-LIB34-018-Q1-E1-E10
US-09-960-352-11486
                                                                                                                                                                                                                                                                                                           US-09-770-791-278/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-960-352-11486
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:

APPLICANT: Wairen, Wesley C.

APPLICANT: Wairen, Wesley C.

APPLICANT: Byatt, John C.

APPLICANT: Byatt, John C.

APPLICANT: Mathialagan, Nagappan

TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION

FILE REFERENCE: 16511.006/37-21(10298)C

CURRENT APPLICATION NUMBER: US/09/960,352

CURRENT FILING DATE: 2001-09-24

NUMBER OF SEQ ID NOS: 15112

SEQ ID NO 11486

LENGTH: 369
                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                         Sequence 278, Application US/09770791 Patent No. US20020062014A1
                                                                                                                                                                             APPLICANT: APPLICANT:
                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 1111

LENGTH: 364

TYPE: DNA

ORGANISM: Bos taurus

OTHER INFORMATION: Clone ID: 05-LIB34-056-Q1-E1-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best
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Patent No. US20020137139A1
                                                                              APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                             APPLICANT: APPLICANT:
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                                                                APPLICANT:
                                                                                                                                                                                                                                          APPLICANT: Gorlach, Jorn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION FILE REFERENCE: 16511.006/37-21(10298)C CURRENT APPLICATION NUMBER: US/09/960,352 CURRENT FILING DATE: 2001-09-24 NUMBER OF SEQ ID NOS: 15112
                    APPLICANT:
                                  APPLICANT:
                                                   APPLICANT:
                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 194 CTGCAAATCCAGTGCT 209
                                                                                                                                                                                                                                                                                                                                                                          21 CTGCAAATCCAGTGCT 36
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                                                                                          Matthew, Abraham V
Ledford, Brooke L.
                                                                                                                                                                                         An, Yong-Qiang
Hamilton, Carol M.
Price, Jennifer L.
             Slader,
                                                             Woessner, Jeffrey P.
Haas, William David
Davis, Keith R.
                            Kricker, Maja
                                            Garcia, Carlos A.
                                                                                                                                                                           Raines, Tracy M.
                                                                                                                               Page,
                                                                                                                                           Yu, Yang
Rameaka, Joshua G.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                          Amy
                                                                                                           Abraham V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.3%;
100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 369;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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RESULT 25
US-10-057-275-3/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: DNA
; ORGANISM: Arabidopsis thaliana
US-09-770-791-278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3, Application US/10057275 Patent No. US20020155545A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 278
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Best Local Similarity
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TITLE OF INVENTION: Expressed Sequences of Arabidopsis
TITLE OF INVENTION: thaliana
FILE REFERENCE: 2029 (PARA-018PRV)
CURRENT APPLICATION NUMBER: US/09/770,791
CURRENT FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/178,480
PRIOR APPLICATION NUMBER: 60/178,480
PRIOR FILING DATE: 2000-01-27
NUMBER OF SEC IN NOC. 2008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 999
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                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       323 AACAACGACTGCTCCT 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                80 AACAACGACTGCTCCT 95
                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0027 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-852-0195
TELEFAX: 415-852-0195
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                      TOPOLOGY: linear
MOLECULE TYPE: CDNA
IMMEDIATE SOURCE:
LIBRARY: Human Pancreas
CLONE: 226152
                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: DOS
SOFTWARE: PRASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/10/057, 275
FILING DATE: 25-Jan-2002
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc
STREET: 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: NEW CHEMOKINES EXPRESSED IN PANCREAS NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Coleman, Roger
Bandman, Olga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16;
                                                                                                                          LENGTH: 402 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/390,740A FILING DATE: February 17, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hoffman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Allen, Keith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.3%; Score 16; DB ilarity 100.0%; Pred. No. 43 Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           U.S.
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; LOCATION: (1)..(405)
; OTHER INFORMATION:
US-10-103-859-1
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                                                                                                                               APPLICANT: Warren, Wesley C.
APPLICANT: Tao, Nengbing
APPLICANT: Tao, Nengbing
APPLICANT: Byatt, John C.
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: MUSCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REPERENCE: 16511.06/37-21(10298)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
SEQ ID NO 3690
LENGTH: 406
                                                                                                                                                                                                                                                                                                                                                                          Sequence 3690, Application US/09960352
Patent No. US20020137139A1
GENERAL INFORMATION:
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CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: US 09/496,273
PRIOR FILING DATE: 2000-02-01
PRIOR APPLICATION NUMBER: US 08/852,212
PRIOR FILING DATE: 1997-05-06
PRIOR FILING DATE: 1996-05-14
PRIOR FILING DATE: 1996-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: White, J.R.
APPLICANT: Pelus, L.M.
TITLE OF INVENTION: Method of Treating Sepsis and ARDS Using Chemokine Beta-9
FILE REFERENCE: 1488.1520003
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 2
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                          TYPE: DNA
ORGANISM: BOS taurus
OTHER INFORMATION: Clone ID: 16-LIB34-030-Q1-E1-D11
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100.0%; Pred. No. 43;
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100.0%;
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100.0%;
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    Score 16;
Pred. No.
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  DB 10;
43;
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                      Length 406;
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LOCATION: (333),(347),(373),(384)
; OTHER INFORMATION: unsure at all n locations
; OTHER INFORMATION: Clone ID: 47-LIB34-011-Q1-E1-D4
US-09-960-352-11005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-960-352-11005
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                                                                                                                                                                                                                                              APPLICANT: Warren, Wesley C.
APPLICANT: Tao, Nengbing
APPLICANT: Tao, Nengbing
APPLICANT: Tao, Nengbing
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REFERENCE: 37-21(10297)C
CURRENT APPLICATION NUMBER: US/99/983,965
CURRENT APPLICATION NUMBER: US/99/983,965
CURRENT FILING DATE: 1990-12-15
PRIOR APPLICATION NUMBER: US 60/465,231
PRIOR APPLICATION NUMBER: US 60/113,678
PRIOR FILING DATE: 1999-12-15
PRIOR FILING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 5912
SEQ ID NO 3483
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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APPLICANT: Mathialagan, Nagappan
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REFERENCE: 16511.006/37-21(10798)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Bos taurus
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                                                              FEATURE:
NAME/KEY: unsure
LOCATION: (76),(81),(87)
OTHER INFORMATION: unsure at all n locations
OTHER INFORMATION: Clone ID: 37-LIB3058-007-Q1-K1-B2
                                                                                                                                                                                  ORGANISM: Bos taurus
                                                                                                                                                                                                           TYPE: DNA
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                                                                                                                                                                                                                                  ENGTH: 414
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100.0%; Pred. No.
4.38;
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Score 16;
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DB 10;
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Length 414;
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US-09-864-761-10902
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                                                                                                                                                                                                                                                                               SOFTWARE: Annomax Sequence Listing Engine vers. SEQ ID NO 10902 LENGTH: 488
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        OTHER INFORMATION: MAP TO AC003103.1

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.9

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.89

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.2

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.6

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.6
                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
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PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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PRIOR FILING DATE: 2000-03-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY FILE REFERENCE: Acomica-x-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                         ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                            TYPE: DNA
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PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT,
FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: PCT/US01/00665
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Chen, Wensheng
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Patent No. USZUCZE PATENT NO. USZUCZE PATENTA PRICAMATION:
APPLICANT: ROSEN et al.
FITTLE OF INVENTION: Nucleic Acids, Proteins, and FILE REFERNCE: PTZ23
CURRENT APPLICATION NUMBER: US/09/764,864
CURRENT FILING DATE: 2001-01-17
                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 32
US-09-764-864-423/c
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                                                                                                                                                                                                SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 423
LENGTH: 615
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 423, Application US/09764864 Patent No. US20020132753A1
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Best Local Similarity
Matches 16; Conserv
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Best Local S
                                                                                                                                                                                                                                                    Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 1792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: The Regents of the University of California APPLICANT: The Government of the United States of America APPLICANT: as represented by the Secretary of the APPLICANT: Department of Health and Human Services TITLE OF INVENTION: SF, a No. US20020051997A1el Family of Taste Receptors FILE REFERENCE: 02307E-098000US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/393,634
CURRENT FILING DATE: 1999-09-10
NUMBER OF SEQ ID NOS: 92
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NAME/KEY: SITE LOCATION: (569)
                                                                                 LOCATION: (513)
OTHER INFORMATION: n equals a,t,g, or
                                 OTHER INFORMATION: n equals a,t,g, or
                                                     NAME/KEY: SITE
LOCATION: (536)
                                                                                                                          FEATURE:
NAME/KEY: SITE
                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Mus sp.
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US20020051997A1
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                                                       (536)
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Ryba, Nick
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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100.0%;
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pred. No. 43;
mismatches
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Pred. No.
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US-09-833-381-94
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 94
LENGTH: 654
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Robison, Keith E.
APPLICANT: Robison, Keith E.
TITLE OF INVENTION: NO. US20020132090Alel Nucleic Acid and Protein Homologs
FILE REFERENCE: 5800-119
CURRENT APPLICATION NUMBER: US/09/833,381
CURRENT FILING DATE: 2001-04-11
PRIOR APPLICATION NUMBER: 09/516,448
PRIOR FILING DATE: 2000-02-29
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Salceda, Susana
APPLICANT: Macina, Roberto
APPLICANT: Recipon, Herve
APPLICANT: Cafferkey, Robert
APPLICANT: Sun, Yongming
APPLICANT: Liu, Chenghua
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                 APPLICANT: Sun, Yongming
APPLICANT: Liu, Chenghua
APPLICANT: Liu, Chenghua
TITLE OF INVENTION: Compositions and Methods Relating to Breast Specific Genes and
FILE REFERENCE: DEX-0269
FILE REFERENCE: DEX-0269
FILE REFERENCE: DEX-0269
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FILE REFERENCE: DEX-0269
FILE REFERENCE: DOM-11-20
FRIOR APPLICATION NUMBER: 60/249,998
FRIOR APPLICATION NUMBER: 60/252,563
FRIOR FILING DATE: 2000-11-22
FRIOR FILING DATE: 2000-11-22
FRIOR FILING DATE: 2000-11-25
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LOCATION: (610)
OTHER INFORMATION: n equals a,t,g, or
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LOCATION: (597)
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LOCATION: (588)
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Pred. No.
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APPLICANT:
APPLICANT:
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RESULT 35
US-09-815-242-4715/c
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CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-03-21
PRIOR PRIOR PRIOR DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-11-27
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
                                                                                                                                                 RESULT 36
US-09-815-242-8625/c
                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Staphylococcus aureus US-09-815-242-4715
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                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 14110
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4715
LENGTH: 1172
TYPE: DNA
Sequence 8625, Application US/09815242
Patent No. US20020061569A1
GENERAL INFORMATION:
APPLICANT: Haselbeck, Robert
APPLICANT: Ohlsen, Kari L.
APPLICANT: Zyskind, Judith W.
APPLICANT: Wall, Daniel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4715, Application US/09815242 Patent No. US20020061569A1
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                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Identification of Essential Genes in TITLE OF INVENTION: Prokaryotes FILE REFERENCE: ELITRA.011A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Haselbeck, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/269,308 PRIOR FILING DATE: 2001-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    311 GGCCCAAGAAAAGGGG 296
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                                                                                                                                                                                                                                        992 AATCCAGTGCTACCAG 977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity nes 16; Conserv
                                                                                                                                                                                                                                                                                                                         Local Similarity 100.0%; nes 16; Conservative
                                                                                                                                                                                                                                                                                   46 AATCCAGTGCTACCAG 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Xu, H. Howard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Carr, Grant J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Trawick, John D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wall, Daniel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yamamoto, Robert T.
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L00.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                        4.38;
                                                                                                                                                                                                                                                                                                                           Score 16; DB; Pred. No. 43
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 9;
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Trawick, John D. Carr, Grant J.

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; LOCATION: (1)...(1179)
US-09-815-242-8625
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PRIOR APPLICATION NUMBER: 60/191,078

PRIOR APPLICATION NUMBER: 60/296,848

PRIOR APPLICATION NUMBER: 60/206,848

PRIOR FILING DATE: 2000-05-23

PRIOR APPLICATION NUMBER: 60/207,727

PRIOR APPLICATION NUMBER: 60/242,578

PRIOR APPLICATION NUMBER: 60/242,578

PRIOR FILING DATE: 2000-10-23

PRIOR APPLICATION NUMBER: 60/253,625

PRIOR APPLICATION NUMBER: 60/257,931

PRIOR APPLICATION NUMBER: 60/257,931

PRIOR FILING DATE: 2000-11-27

PRIOR FILING DATE: 2000-11-27

PRIOR FILING DATE: 2000-11-27

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PRIOR FILING DATE: 2000-11-27

PRIOR FILING DATE: 2000-11-27

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PRIOR FILING DATE: 2000-11-27

PRIOR FILING DATE: 2000-11-27

PRIOR FILING DATE: 2000-12-26

PRIOR FILING DATE: 2000-13-27
                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 222
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 34
LENGTH: 1549
                                                                                                   Matches
                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 34, Application US/09813358 Patent No. US20020048759A1
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SEQ ID NO 8625
LENGTH: 1179
                                                                                                                                Best
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Xu, Jiangchun
APPLICANT: Pyle, Ruth
APPLICANT: Stolk, John A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN AND ENDOMETRIAL CANCER
FILE REFERENCE: 210121.501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/813,358 CURRENT FILING DATE: 2001-03-21
                                                                                                                                                                                                            FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(1549)
OTHER INFORMATION: n = A,T,C or G
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PRIOR FILING DATE: 2001-02-16
NUMBER OF SEQ ID NOS: 14110
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APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/815,242 CURRENT FILING DATE: 2001-03-21
                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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                                               277 CTGCTGCAACACCCCT 292
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  57
                                                                                                                        Local Similarity
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CTGCTGCAACACCCCT 72
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                                                                                                 Conservative
                                                                                                                        4.3%;
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                                                                                                                      Score 16;
Pred. No.
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                                                                                                 Mismatches
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                                                                                                                                             DB 10; Length 1549;
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CURRENT APPLICATION NUMBER: US/09/801,368
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US 09/487,558
PRIOR FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: US 60/160,587
PRIOR APPLICATION NUMBER: US 60/160,587
PRIOR FILING DATE: 1999-10-20
NUMBER OF SEQ ID NOS: 440
SOFTWARE: Patentin version 3.0
SEQ ID NO 41
LENGTH: 1989
TYPE: DNA
ORGANISM: Aspergillus nidulans
US-09-801-368-41
                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-822-849A-343/c
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                                                CURRENT APPLICATION NUMBER: US/09/822,849A
CURRENT FILING DATE: 2001-09-04
PRIOR APPLICATION NUMBER: 60/195,582
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 598
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 343
FENCEN: 2062
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Best Local
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APPLICANT:
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APPLICANT: Cali,
APPLICANT: Hecht
                                                                                                                                                                                   APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING NOVEL SECRETED PROTEINS
FILE REFERENCE: GIN 6403
                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Summers, Eric TITLE OF INVENTION: Methods for Improving Secondary Metabolite Production in Fungi-
FILE REFERENCE: 109272.147
                                                                                                                                                                                                                                           APPLICANT:
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APPLICANT:
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ORGANISM: Homo sapiens
                                    LENGTH: 2053
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                                                                                                                                                                                                                                       Gulukota, Kamalakar
Graham, James R
                                                                                                                                                                                                                                                                              Resnick, Richard J.
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Silva, Jeff
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Salama, Sofie
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No. US20020128250Alman, Thea
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                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/09822849A
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US-09-822-849A-343

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RESULT 40
US-09-802-669-94/c
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; LOCATION: (1782)...(1813)
US-09-802-669-94
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-922-261-359
; LENGTH: 2702
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-922-261-359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 94, Application US/09802669 Patent No. US20020004490A1
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                                                                                PRIOR APPLICATION NUMBER: US/09/461,697
PRIOR FILING DATE: 1999-12-14
NUMBER OF SEQ ID NOS: 466
SOFTWARE: FAST-SEQ for Windows Version 4.0
SEQ ID NO 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best
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APPLICANT: Zhang, Hong
TITLE OF INVENTION: Antisense Compound Modulation of Fas Mediated Signaling
FILE REFERENCE: ISPH-545
CURRENT APPLICATION NUMBER: US/09/802,669
CURRENT FILING DATE: 2001-03-09
PRIOR APPLICATION NUMBER: US 09/665,615
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: US 09/290,640
PRIOR APPLICATION NUMBER: US 09/290,640
PRIOR FILING DATE: 1999-04-12
NUMBER OF SEC ID NOS: 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Dean, Nicholas M. APPLICANT: Marcusson, Eric G.
                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                 APPLICANT: PORTBURY, STUART D.
APPLICANT: Puranam, Kasturi
APPLICANT: Katz, Lawrence C.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING
TITLE OF INVENTION: AND TREATING CONDITIONS, DISORDERS, OR DISEASES INVOLVING
TITLE OF INVENTION: CELL DEATH
FILE REFERENCE: 10001-005-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/922,261
CURRENT FILING DATE: 2001-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: COGENT NEUROSCIENCE, Inc. APPLICANT: Lo, Donald C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 2165
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1096 CAAAGTGCCGGGATCA 1081
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            306 CAAGGCCCAAGAAAAG 321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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Thomas, Mary Beth
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В
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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1156
LENCTH: 3294
TYPE: DNA
ORGANISM: Homo sapiens
US-09-764-860-1156
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US-09-834-291-1/c
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TITLE OF INVENTION: p53 Binding Areas
FILE REFERENCE: 4121-112
FURRENT APPLICATION NUMBER: US/09/834,291
CURRENT FILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: DCT/DE99/03343
PRIOR FILING DATE: 1999-10-18
PRIOR APPLICATION NUMBER: DE 198 47 779.1
PRIOR APPLICATION NUMBER: D2 198 47 779.1
PRIOR FILING DATE: 1998-10-16
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APPLICANT: Krammer, Peter
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
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Best Local Similarity
                                                                Query Match
Best Local Similarity
"atches 16; Conser"
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                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/764,860 CURRENT FILING DATE: 2001-01-17
                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC008
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TYPE: DNA
ORGANISM: Homo Sapiens
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819 GGTACCAGTCCTTCTG 834
                                       228 GGTACCAGTCCTTCTG 243
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100.0%; Pred. No. 44;
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RESULT 44 US-09-764-877-2431

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RESULT 45

US-09-764-869-2170

Sequence 2170, Application US/09764869

Patent No. US20020061521A1

GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC007

CURRENT APPLICATION UMBER: US/09/764,869

CURRENT APPLICATION UMBER: US/09/764,869

CURRENT APPLICATION OF ENDINGLED OF SEQ ID NOS: 2442

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 2170

LENGTH: 17581

TYDE: DNA
ORGANISM: Homo sapiens
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APPLICANT: Rosen et al.

TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies

FILE REFERENCE: PC005

CURRENT APPLICATION NUMBER: US/09/764,877

CURRENT FILING DATE: 2001-01-17

Prior application data removed - refer to PALM or file wrapper NUMBER: OF SEQ ID NOS: 4031

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 2431

LENGTH: 11749

TYPE: DNA

ORGANISM: Homo sapiens

US-09-764-877-2431
Search completed: November 8, 2002, 02:05:12 Job time : 41.0364 secs
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                                                                                   2661 AAGAAAAGGGGAAGTT 2676
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Total number of hits satisfying chosen parameters:
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Maximum DB seq length: 200000000
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369
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/cgn2_6/ptodata/1/pna/US6006_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.

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1524	1524	1524	1524	1524	1524	1524	1524	488	488	488	488	414	483	483	369	369	369	369	369	369	Match Length
42	42	42	42	42	42	42	40	34	34	16	16	42	42	29	42	36	31	28	25	٣	BG
US-10-216-166-15	US-10-216-165-15	US-10-216-164-15	US-10-216-163-15	US-10-216-162-15	US-10-216-160-15	US-10-216-159A-15	US-10-119-480-15	US-09-909-627-6100	US-09-904-809-18421	US-09-277-227-6100	US-09-234-611-18421	US-10-212-677-283	US-10-235-954-340	US-09-757-034-340	US-10-212-677-199	US-09-970-966-199	US-09-825-294-199	US-09-713-550-199	US-09-656-668-199	PCT-US01-45395-199	ID
Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl		Sequence 18421, A		18421	283,	340,	340,		199,	Sequence 199, App	Sequence 199, App	Sequence 199, App	Sequence 199, App	Description

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RESULT 1
PCT-US01-45395-199
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NAME/KEY: misc_feature
LOCATION: 29, 345
OTHER INFORMATION: n =
PCT-US01-45395-199
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 199, Application PC/TUS0145395

GENERAL INFORMATION:
APPLICANT: Corixa Corporation
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Filing, Steven P.
APPLICANT: Filing, Steven P.
APPLICANT: Molesh, David Alan
APPLICANT: Molesh, David Alan
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
TITLE REFERENCE: 210121.48401PC
                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
LENGTH: 369
                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                 CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTGTCTGCCGGGTACCAGTCCTT
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US-10-218-631-15
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US-10-218-784-15
US-10-218-930-15
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APPLICANT: Xu, Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

TITLE OF INVENTION: OVARIAN TUMOR SEQUENCES

TITLE OF INVENTION: METHODS OF USE THEREFOR

FILE REFERENCE: 210121.484C3

FILE REFERENCE: 210121.484C3

CURRENT APPLICATION UNMBER: US/09/656,668

CURRENT FILING DATE: 2000-09-07

NUMBER OF SEQ ID NOS: 199
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OF USE THEREFOR
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Pred. No. 1.1e-188;
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Sequence 199, Application US/09825294

GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Filing, Steven P.

APPLICANT: Filing, Steven P.

TITLE OF INVENTION: THERAPY AND METHODS FOR THE TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 199

LENGTH: 369
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CURRENT APPLICATION NUMBER: US/09/713,550
CURRENT FILING DATE: 2000-11-14
NUMBER OF SEQ ID NOS: 205
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 199
LENGTH: 369
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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; LCCATION: (1)...(369)
; OTHER INFORMATION: n = A,T,C or
US-09-713-550-199
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Best Local Similarity
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Pred. No. 1.1e-188;
; Mismatches 0;
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RESULT 5
US-09-970-966-199
; Sequence 199, Application US/09970966
; GENERAL INFORMATION:
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                                                                                                         ; FEATURE:

, NAME/KEY: misc_feature

; LOCATION: 29, 345

; OTHER INFORMATION: n =

US-09-970-966-199
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                                                                                                                                                                                    TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE RESERVE: 21011.484C6
CURRENT APPLICATION NUMBER: US/09/970,966
CURRENT FILING DATE: 2001-10-02
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
LENGTH: 369
TYPE: DNA
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Matches
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Fling, Steven P.
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Pred. No. 1.1e-188;
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                                                   Score 369; DB 36;
Pred. No. 1.1e-188;
Mismatches 0;
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CURRENT FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 288
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
LENGTH: 369
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APPLICANT: MCNeill, Patricia D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C7
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                                               CCGCAAGTCCTGTGCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
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; NAME/KEY: SITE
; LOCATION: (443)
; OTHER INFORMATION: n e
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NUMBER OF SEQ ID NOS: 1440
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 340
                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                             Matches 317;
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CURRENT APPLICATION NUMBER: US/09/757,034
CURRENT FILING DATE: 2001-01-09
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NAME/KEY: SITE
LOCATION: (435)
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NAME/KEY: SITE
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(367)
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                                                                                                                                                                                                                                                                                                                         Score 317; DB 29;
Pred. No. 1.7e-160;
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RESULT 8 US-10-235-954-340

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PRIOR APPLICATION NUMBER: 09/757,034
PRIOR FILING DATE: 2001-01-09
PRIOR APPLICATION NUMBER: 60/179,065
PRIOR FILING DATE: 2000-01-31
PRIOR PPLICATION NUMBER: 60/180,628
PRIOR FILING DATE: 2000-02-04
NUMBER OF SEQ ID NOS: 1440
SEQ ID NO 340
LENGTH: 483
TYPE: DNA
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                                                                                                                                Sequence 283, Application US/10212677 GENERAL INFORMATION:
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CURRENT FILING DATE: 2002-09-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: PM002C1N
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TITLE OF INVENTION: Nucleic Acids, Proteins,
APPLICANT: Chenault, Ruth A.
APPLICANT: Xu, Jiangchun
APPLICANT: Fanger, Gary R.
APPLICANT: Harlocker, Susan L.
APPLICANT: MCNeill, Patricia D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature
LOCATION: (443)
OTHER INFORMATION: n equals a,t,g,
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OTHER INFORMATION: n equals
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LOCATION: (435)
OTHER INFORMATION: n equals
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100.0%;
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 Mismatches

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; LENGTH: 414
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-212-677-283
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                                                                                                                                           ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(488)
OTHER INFORMATION: n = A,T,C or
US-09-234-611-18421
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US-09-234-611-18421
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CURRENT APPLICATION NUMBER: US/10/212,677
CURRENT FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 288
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                         Query Match
Best Local S
                                                                                               Matches 315;
                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                                             LENGTH: 488
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                                             CTGCCTCGGCCCTCA 352
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                                                                                                          Similarity
                                                                                               Conservative
                                                                                               85.4%; Score 315; DB 16;
100.0%; Pred. No. 2e-159;
tive 0; Mismatches 0;
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                                                                                                                       Length 488;
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RESULT 12
US-09-904-809-18421
Sequence 18421, Application US/09904809
GENERAL INFORMATION:
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: LOCATION: (1)...(488)
: OTHER INFORMATION: n = A,T,C
US-09-277-227-6100
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US-09-277-227-6100
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TITLE OF INVENTION: ROVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS cDNA LIBRARIES
FILE REFERENCE: 20411-766
CURRENT APPLICATION NUMBER: US/09/277,227
CURRENT FILING DATE: 1999-03-25
NUMBER OF SEQ ID NOS: 23680
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 6100
SEQ ID NO 6100
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GENERAL INFORMATION
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                                                                                                                                                                                                                                    AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCATCAGCGGCCT 209
                                                                                                                                                                        GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT
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100.0%;
                                                                                                                                                                                                                                                                                                                                                                                       Score 315; Pred. No.
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APPLICANT: Hyseq, Inc.
APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS cDNA LIBRARIES
FILE REFERENCE: 20411-766
CURRENT APPLICATION NUMBER: US/09/909,627
CURRENT FILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: 09/277,227
PRIOR APPLICATION NUMBER: 09/277,227
PRIOR FILING DATE: 1999-03-23
NUMBER OF SEQ ID NOS: 23680
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 6100
; OTHER INFORMATION: n = US-09-909-627-6100
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; GENERAL INFORMATION:
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CURRENT FILING DATE: 2001-07-12
PRIOR APPLICATION NUMBER: 09/234,611
PRIOR FILING DATE: 1999-01-22
NUMBER OF SEQ ID NGS: 21025
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 18421
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Best Local
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                                     NAME/KEY: misc_feature LOCATION: (1)...(488)
                                                                            FEATURE:
                                                                                                ORGANISM: Homo sapiens
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TITLE OF INVENTION: HOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-757
                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 488
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                              LENGTH: 488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        379 CTGCCTCGGCCCTCA 393
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LENGTH: 1524
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Best Local
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                                                                                                                                                                                                                                        Query Match
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CURRENT FILING DATE: 2002-04-09
NUMBER OF SEQ ID NOS: 246
Prior Application removed - See File Wrapper or Palm
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APPLICANT: Desnoyers, Luc
APPLICANT: Goddard, Audre
APPLICANT: Godowski, Paul
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                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: P3530P1C1
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Grimaldi, J. Christopher
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; SEQ ID NO 15
; LENGTH: 1524
; TYPE: DNA
; ORGANISM: Homo Sapien
US-10-216-159A-15
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CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-01-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR APPLICATION NUMBER: 60/078973
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                                                                                                                                                                                                                                                                                                                                                                                                       Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 246
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150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
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                                        GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
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Grimaldi, J. Christopher
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Gerritsen, Mary
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100.0%; Pred. No. 2e-159;
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APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
ITILE OF INVENTION: SECETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C8
CURRENT APPLICATION NUMBER: US/10/216,160
CURRENT FILING DATE: 2002-08-09
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
TYPE: DNA
ORGANISM: Homo Sapien
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Best Local Similarity 100.
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APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, J. Chri
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                                                                330 CTGCCTCGGCCCTCA 344
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                                  CTGCCTCGGCCCTCA 484
                                                                                               GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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330 CTGCCTCGGCCCTCA 344
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
TYPE: DNA
ORGANISM: Homo Sapien
US-10-216-162-15
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PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/063349
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/064873
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/079910
PRIOR APPLICATION NUMBER: 60/079294
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PRIOR APPLICATION NUMBER: 60/079294
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PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079656
PRIOR FILING DATE: 1998-03-26
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Best Local Similarity
Matches 315; Conserv
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APPLICANT:
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CURRENT FILING DATE: 2002-08-09
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APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C2
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APPLICANT:
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                                                                                                 GCATCAGCTGCTACACCCCCTCTTTGTAACGGGCCCAAGGCCCAAGAAAAGGGGAAGTT 329
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Gerritsen, Mary
Goddard, Audrey
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Grimaldi, J. Christopher
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; ORGANISM: Homo Sapien
US-10-216-163-15
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CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
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270 GCATCAGCTGCTAACACCCCTCTTTGTAACGGGCCCAAGGCCCAAGAAAAGGGGAAGTT 329
                                                                                                                                                                                                                                                                      290 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
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FILING DATE: 1998-03-26
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FILING DATE: 1998-03-25
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FILING DATE: 1997-12-17
                                      GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 269
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Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
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Smith, Victoria
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                                                                                                                                                                                                                                                                                                                                                   85.4%; Score 315; DB 42; Length 1524; 100.0%; Pred. No. 2e-159; tive 0; Mismatches 0; Indels 0
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; TYPE: DNA
; ORGANISM: Homo Sapien
US-10-216-164-15
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CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/05287
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/063549
PRIOR PILING DATE: 1997-10-18
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
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LENGTH: 1524
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PRIOR FILING DATE: 1998-03-27
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APPLICANT: Desnoy
APPLICANT: Gerrit
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PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
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                                      290 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 349
210 GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 269
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                                                                                                                                                                                          90 GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
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FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 1998-03-26
                                                            AAGTGATGGAGCAAAGTGCCGGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
                                                                                                                 GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 289
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                                                                                                                                                                                                                                                                                               Similarity
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Goddard, Audrey
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                                                                                                                                                                                                                                                                     85.4%; Score 315; DB 42; Length 1524; 100.0%; Pred. No. 2e-159; tive 0; Mismatches 0; Indels 0
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CTGCCTCGGCCCTCA 344

US-10-216-165-15

GENERAL INFORMATION: Sequence 15,

Application US/10216165

APPLICANT: Baker, Kevin P.
APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mar
APPLICANT: Goddard, Audre

APPLICANT:

Godowski, Paul J. Grimaldi, J. Christopher

Goddard, Audrey

APPLICANT: APPLICANT:

Gurney, Austin L. Smith, Victoria

APPLICANT:

APPLICANT:

Watanabe, Colin L. Wood, William I.

Stephan, Jean-Philippe F.

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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                              Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 246
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                                                                                                          230 GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
                                                       90 GCTCCTCCCCGGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
                                                                                                                                          30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
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PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
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                                                                                                                                                                                                                                             Remaining Prior Application
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
                                                                                                                       Best Local Similarity
Matches 315; Conserv
                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3530P1C9
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
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CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
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                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1998-03-20
                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1998-03-27
                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/079728
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                                                                                                                                                                                                                                    TYPE: DNA
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                                               90 GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
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                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 1998-03-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Gerritsen, Mary
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                                                                                                                         ilarity 100.0%;
Conservative (
                                                                                                                                                               85.48;
                                                                                                                                                                                                                                                                                                              data removed - See File Wrapper or
                                                                                                                           0;
                                                                                                                                            Score 315; Pred. No.
                                                                                                                             Mismatches
                                                                                                                                              DB 42;
2e-159;
                                                                                                                               0;
                                                                                                                                                               Length 1524;
                                                                                                                                 Indels
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PRIOR APPLICATION NUMBER: 10/119,480 PRIOR FILING DATE: 2002-04-09 PRIOR PPLICATION NUMBER: 60/059113 PRIOR FILING DATE: 1997-09-17 PRIOR PPLICATION NUMBER: 60/062287 PRIOR PPLICATION NUMBER: 60/062287 PRIOR FILING DATE: 1997-10-17

PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25

PRIOR APPLICATION NUMBER: 60/079656

FILING DATE: 1998-03-26

PRIOR APPLICATION NUMBER:

60/078910

PRIOR APPLICATION NUMBER: 60/079728 PRIOR FILING DATE: 1998-03-27

PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17

FILE REFERENCE: P3530P1C7
CURRENT APPLICATION NUMBER: US/10/216,165
CURRENT FILING DATE: 2002-08-09

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US-10-216-165-15

ORGANISM: Homo Sapien

TYPE: DNA LENGTH:

1524

Query Match Best Local Matches

Similarity

85.4%;

Score 315; Pred. No. Mismatches

315;

Conservative

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Indels

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RESULT 22
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                                                                                                  ; TYPE: DNA
; ORGANISM: Homo Sapien
US-10-216-167-15
                                Query Match
Best Local Similarity
Matches 315; Conserv
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LENGTH: 1524
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CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/62287
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-17
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30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
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FILING DATE: 1998-03-20
APPLICATION NUMBER: 60/079294
FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
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FILING DATE: 1997-12-17
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Grimaldi, J. Christopher
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Gerritsen, Mary
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                                  Conservative
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                                                85.4%;
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                                Score 315; DB; Pred. No. 2e-
0; Mismatches
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. 2e-159;
ches 0;
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                                                                Length 1524;
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; ORGANISM: Hor
US-10-216-168-15
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PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/063287
PRIOR PILING DATE: 1997-10-17
PRIOR PILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
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PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079656
PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-26
PRIOR FILING DATE: 1998-03-27
PRIOR FILING DATE: 1998-03-27
                                                                                                             NUMBER OF SEQ ID SEQ ID NO 15
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Query Match
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CURRENT FILING DATE: 2002-08-09
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                                                                                                                                                  Remaining Prior Application data removed -
                                                                                           LENGTH: 1524
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                                                      Homo
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Smith, Victoria
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Grimaldi, J. Christopher
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Gerritsen, Mary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wood, William I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stephan, Jean-Philippe F. Watanabe, Colin L.
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                                                      Sapien
85.4%;
Score
315;
DВ
                                                                                                                                                  See File Wrapper or PALM.
42;
Length 1524;
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CURRENT APPLICATION NUMBER: US/10/218,612
CURRENT FILING DATE: 2002-08-12
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/062287
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                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
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GENERAL I
                                    SEQ ID NO 15
LENGTH: 15
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: WOOG, WIlliam I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
                                                                        Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 246
                                                                                                              PRIOR FILING DATE: 1998-03-27
                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/063549 PRIOR FILING DATE: 1997-10-28
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                                                                                                                                                                         APPLICATION NUMBER: 60/079656
                                                                                                                                  APPLICATION NUMBER: 60/079728
                                                                                                                                                       FILING DATE:
                                                                                                                                                                                             FILING DATE: 1998-03-25
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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                                                                                                                                                       1998-03-26
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PRIOR APPLICATION.

PRIOR FILING DATE: 1998-03-25

PRIOR FILING DATE: 1998-03-26

PRIOR APPLICATION NUMBER: 60/079728

PRIOR FILING DATE: 1998-03-27

PRIOR FILING DATE: 1998-03-27
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                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/218,631
CURRENT FILING DATE: 2002-08-12
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/05287
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3530P1C14
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                                                                                                        APPLICATION NUMBER: 60/078910 FILING DATE: 1998-03-20 APPLICATION NUMBER: 60/079294 FILING DATE: 1998-03-25
                                                                                                                                                                                         APPLICATION NUMBER: 60/064103
FILING DATE: 1997-10-31
APPLICATION NUMBER: 60/069873
FILING DATE: 1997-12-17
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                  Prior Application data removed -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stephan, Jean-Philippe Watanabe, Colin L. Wood, William I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gurney, Austin L. Smith, Victoria
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Pred. No.
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thes 0;
                See File Wrapper or PALM
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                                            CURRENT FILING DATE: US/10/218,765
CURRENT FILING DATE: 2002-08-12
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/064103
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PRIOR APPLICATION NUMBER: 60/069873
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LENGTH: 1524
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Matches 315; Conserv
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APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3550P1C19
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Gerritsen, Mary
Goddard, Audrey
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OR FILING DATE: 1998-09-10
OR APPLICATION NUMBER: 60/099812
OR FILING DATE: 1998-09-10
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OR APPLICATION NUMBER: 60/090695
OR FILING DATE: 1998-06-25
OR APPLICATION NUMBER: 60/091982
OR FILING DATE: 1998-07-07
OR APPLICATION NUMBER: 60/095302
OR FILING DATE: 1998-08-04
OR APPLICATION NUMBER: 60/095318
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PR APPLICATION NUMBER: 60/099596

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PR APPLICATION NUMBER: 60/099598

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PR APPLICATION NUMBER: 60/099803

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APPLICATION NUMBER: 60/
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APPLICATION NUMBER: 60/095916
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APPLICATION NUMBER: 60/096791
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APPLICATION NUMBER: 60/100627
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DR APPLICATION NUMBER: 60/130232
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OR FILING DATE: 1998-11-17
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Best Local Similarity
                                                                                            APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C18
CURRENT APPLICATION NUMBER: US/10/218,784 CURRENT FILING DATE: 2002-08-12 PRIOR APPLICATION NUMBER: 10/119,480 PRIOR FILING DATE: 2002-04-09 PRIOR APPLICATION NUMBER: 60/059113 PRIOR FILING DATE: 1997-09-17
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OR FILING DATE: 1999-08-17
OR APPLICATION NUMBER: 60/151733
OR FILING DATE: 1999-08-31
OR APPLICATION NUMBER: 60/164418
OR FILING DATE: 1999-11-09
OR APPLICATION NUMBER: 60/166361
OR FILING DATE: 1999-11-16
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APPLICATION NUMBER: 60/146963
FILING DATE: 1999-08-03
APPLICATION NUMBER: 60/149320
FILING DATE: 1999-08-17
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FILING DATE: 1999-12-07
APPLICATION NUMBER: 60/169835
FILING DATE: 1999-12-07
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Gerritsen, Mary
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Φ.			7	NUMBER	NUMBER: 60	1999-04-21	1999-04	NUMBER: 60	NUMBER: 60/126//3	1999-03-23	NUMBER: 6	1999-03-19	1999-03	NUMBER: 6	1999-02-10	NIIMBER.	NUMBER: 6	1999-01-12	1999-01-12	NUMBER: 60	NUMBER	: 1998-12-23	NUMBER: 60	NUMBER: 60/113296	3	ດ້	NUMBER: 60/108849	: 1998-11-17	1998-11-1/ NUMBER: 60/108801	NUMBER: 60/108787	1998-11-03	1998-10-30	1998-10-29 NIMBER: 60/106464	60	NUMBER: 60/106178	09-24	1998-09-24 NIMBER: 60/101922	NUMBER: 60/101916	1998-09-25	1998-09-24	R: 6	1998-09-24	1998-09-23	H	8-09-17	1998-09-18	BER: 60	NUMBER: 60/100627 1998-09-16	9-15	1998-09-15 NUMBER: 60/100390	ER: 60	1998-09-11

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OR APPLICATION NUMBER: 60/062287
OR REPLING DATE: 1997-10-28
OR APPLICATION NUMBER: 60/06349
OR FILING DATE: 1997-10-31
OR APPLICATION NUMBER: 60/079294
OR FILING DATE: 1997-12-17
OR APPLICATION NUMBER: 60/079294
OR FILING DATE: 1998-03-25
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OR APPLICATION NUMBER: 60/079296
OR FILING DATE: 1998-03-27
OR APPLICATION NUMBER: 60/081919
OR FILING DATE: 1998-03-27
OR APPLICATION NUMBER: 60/081919
OR FILING DATE: 1998-04-15
OR APPLICATION NUMBER: 60/082804
OR FILING DATE: 1998-04-15
OR APPLICATION NUMBER: 60/082804
OR FILING DATE: 1998-05-06
OR APPLICATION NUMBER: 60/085323
OR FILING DATE: 1998-05-13
OR APPLICATION NUMBER: 60/085323
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OR APPLICATION NUMBER: 60/080557
OR APPLICATION NUMBER: 60/080532
OR FILING DATE: 1998-06-17
OR APPLICATION NUMBER: 60/080532
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OR APPLICATION NUMBER: 60/090472
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OR APPLICATION NUMBER: 60/09

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; Sequence 15, Application US/10218849
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Gurney, Austin L.
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US-10-218-849-15
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OR FILING DATE: 1999-08-03

OR APPLICATION NUMBER: 60/149320

OR FILING DATE: 1999-08-17

OR APPLICATION NUMBER: 60/149638

OR FILING DATE: 1999-08-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/169445
FILING DATE: 1999-12-07
APPLICATION NUMBER: 60/169495
FILING DATE: 1999-12-07
APPLICATION NUMBER: 60/169835
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FILING DATE: 1999-11-16
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FILING DATE: 1999-07-28
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APPLICATION NUMBER: 60/141037
FILING DATE: 1999-06-23
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                                                                                                                                                                                                                                                                             GCATCAGCTGCTACACCCCCTCTTTGTAACGGGCCCAAGAAAAAGGGGGAAGTT
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             Desnoyers, Luc
Gerritsen, Mary
Goddard, Audrey
Godowski, Paul J.
Grimaldi, J. Christopher
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; ORGANISM: Homo Sapien US-10-218-930-15
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US-10-218-930-15
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                                  CURRENT APPLICATION NUMBER: US/10/218,930
CURRENT FILING DATE: 2002-08-12
Prior Application removed - See File Wrap;
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
TYPE: DNA
                                                                                                                                                                                                                                                                                                           APPLICANT: Baker, Kevin P
APPLICANT: Desnoyers, Lu
APPLICANT: Geritsen, Ma
APPLICANT: Goddard, Audra
APPLICANT: Godowski, Pau,
APPLICANT: Grimaldi, J. o
APPLICANT: Gurney, Austin
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LENGTH: 1524
TYPE: DNA
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Matches
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                                                                                                                                                                                                                                APPLICANT: Stephan, Jean-Philippe F. APPLICANT: Watanabe, Colin L. APPLICANT: Wood, William I.
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APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C11
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CURRENT FILING DATE: 2002-08-12
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NUMBER OF SEQ ID NOS: 246
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Gerritsen, Mary
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Grimaldi, J. Christopher
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PRIOR FILING DATE: 2002-04-09
PRIOR PELICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/064103
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Best Local Similarity
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CURRENT FILING DATE: 2002-08-12
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             OR FILING DATE: 1998-03-25

OR APPLICATION NUMBER: 60/079656

OR FILING DATE: 1998-03-26

OR APPLICATION NUMBER: 60/079728

OR FILING DATE: 1998-03-27

OR FILING DATE: 1998-04-15

OR FILING DATE: 1998-04-15
                                                                                                                                                                                 APPLICATION NUMBER: 60/
FILING DATE: 1998-03-20
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APPLICATION NUMBER: 60/069873
FILING DATE: 1997-12-17
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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DR APPLICATION NUMBER: 60/095318
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DR APPLICATION NUMBER: 60/095916
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DR APPLICATION NUMBER: 60/131022

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DR APPLICATION NUMBER: 60/131291

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OR APPLICATION NUMBER: 60/123618
OR FILING DATE: 1999-03-10
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APPLICATION NUMBER: 60/146222
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APPLICATION NUMBER: 60/144758
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FILE REFERENCE: P3530PICI2

CURRENT APPLICATION NUMBER: US/10/219,003

CURRENT FILING DATE: 2002-08-12

PRIOR APPLICATION NUMBER: 10/119,480

PRIOR FILING DATE: 2002-04-09

PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/05287

PRIOR APPLICATION NUMBER: 60/062287

PRIOR FILING DATE: 1997-10-17

PRIOR FILING DATE: 1997-10-17

PRIOR APPLICATION NUMBER: 60/063549

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PRIOR APPLICATION NUMBER: 60/063549

PRIOR FILING DATE: 1997-10-28

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PRIOR APPLICATION NUMBER: 60/151733
PRIOR TILING DATE: 1999-08-31
PRIOR APPLICATION NUMBER: 60/164418
PRIOR APPLICATION NUMBER: 60/16361
PRIOR FILING DATE: 1999-11-16
PRIOR FILING DATE: 1999-11-16
PRIOR APPLICATION NUMBER: 60/169445
PRIOR APPLICATION NUMBER: 60/169445
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APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Goddwski, Paul J
APPLICANT: Grimmaldi, J. Chr
APPLICANT: Gurney, Austin I
                                                                                                                                                                                                                                                       APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE DEFINITION: ACIDS ENCODING THE SAME
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315; Conserv
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Godowski, Paul J.
Gorimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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; Pred. No. 2e-159;
0; Mismatches 0;
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PRIOR FILING DATE: 1998-09-10 PRIOR PPLICATION NUMBER: 60/099812 PRIOR FILING DATE: 1998-09-10 PRIOR APPLICATION NUMBER: 60/099816 PRIOR FILING DATE: 1998-09-10 PRIOR APPLICATION NUMBER: 60/100038 PRIOR FILING DATE: 1998-09-11 PRIOR APPLICATION NUMBER: 60/100385 PRIOR FILING DATE: 1998-09-15 PRIOR PRIOR APPLICATION NUMBER: 60/100390 PRIOR FILING DATE: 1998-09-15 PRIOR APPLICATION NUMBER: 60/100390 PRIOR APPLICATION NUMBER: 60/100627	A APPLICATION NUMBER: 60 A FILING DATE: 1998-08-1 A APPLICATION NUMBER: 60 A FILING DATE: 1998-08-2 A APPLICATION NUMBER: 60 A FILING DATE: 1998-08-3 A APPLICATION NUMBER: 60 A FILING DATE: 1998-09-0 A APPLICATION NUMBER: 60 A APPLICATION NUMBER: 60 A APPLICATION NUMBER: 60 A APPLICATION NUMBER: 60 A APPLICATION NUMBER: 60 A APPLICATION NUMBER: 60 A APPLICATION NUMBER: 60 A APPLICATION NUMBER: 60 A APPLICATION NUMBER: 60	APPLICATION NUMBER: 6 FILING DATE: 1998-06- FILING DATE: 1998-06- FILING DATE: 1998-06- APPLICATION NUMBER: 6 FILING DATE: 1998-06- APPLICATION NUMBER: 6 FILING DATE: 1998-07- FILING DATE: 1998-07- FILING DATE: 1998-08- FILING DATE: 1998-08- APPLICATION NUMBER: 6 FILING DATE: 1998-08- APPLICATION NUMBER: 6 APPLICATION NUMBER: 6 APPLICATION NUMBER: 6 APPLICATION NUMBER: 6 APPLICATION NUMBER: 6 FILING DATE: 1998-08- APPLICATION NUMBER: 6 FILING DATE: 1998-08- FILING DATE: 1998-08-	APPLICATION NUMBER: 60 APPLICATION NUMBER: 60 APPLICATION NUMBER: 60 APPLICATION NUMBER: 60 FILING DATE: 1998-05-1 APPLICATION NUMBER: 60 FILING DATE: 1998-05-2 APPLICATION NUMBER: 60-1 APPLICATION NUMBER: 60-1 APPLICATION NUMBER: 60-1 FILING DATE: 1998-06-1 APPLICATION NUMBER: 60 FILING DATE: 1998-06-1 APPLICATION NUMBER: 60-6-1 APPLICATION NUMBER: 60-6-1 APPLICATION NUMBER: 60-6-1 APPLICATION NUMBER: 60-6-1 APPLICATION NUMBER: 60-6-1 APPLICATION NUMBER: 60-6-1	PRIOR APPLICATION NUMBER: 60/069873  PRIOR APPLICATION NUMBER: 60/078910  PRIOR APPLICATION NUMBER: 60/079910  PRIOR FILING DATE: 1998-03-20  PRIOR FILING DATE: 1998-03-25  PRIOR APPLICATION NUMBER: 60/079294  PRIOR FILING DATE: 1998-03-25  PRIOR APPLICATION NUMBER: 60/079728  PRIOR APPLICATION NUMBER: 60/079728  PRIOR FILING DATE: 1998-03-27  PRIOR APPLICATION NUMBER: 60/081819  PRIOR APPLICATION NUMBER: 60/081955  PRIOR APPLICATION NUMBER: 60/081955  PRIOR FILING DATE: 1998-04-15  PRIOR FILING DATE: 1998-04-15  PRIOR FILING DATE: 1998-04-15  PRIOR FILING DATE: 1998-04-28  PRIOR FILING DATE: 1998-04-15  PRIOR APPLICATION NUMBER: 60/082804  PRIOR FILING DATE: 1998-04-28
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RESULT 32
US-10-219-010-15
Sequence 15, Applicati
GENERAL INFORMATION:
GENERAL INFORMATION:
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APPLICANT: Baker, Key
APPLICANT: DESNOYER
APPLICANT: Goddard,
APPLICANT: Goddwski
APPLICANT: Godowski
APPLICANT: Grimaldi
APPLICANT: Gurney,
APPLICANT: Smith, V.
APPLICANT: Stephan,
APPLICANT: Watanabe,
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OR APPLICATION NUMBER: 60/144758
OR FILING DATE: 1999-07-26
OR APPLICATION NUMBER: 60/145598
OR FILING DATE: 1999-07-26
OR FILING DATE: 1999-07-28
OR APPLICATION NUMBER: 60/146963
OR APPLICATION NUMBER: 60/146963
OR APPLICATION NUMBER: 60/149320
OR FILING DATE: 1999-08-01
OR APPLICATION NUMBER: 60/151733
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 INVENTION:
                   INVENTION:
                                                           Desnoyers, Luc
Gerritsen, Mary
Goddard, Audrey
Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
Stephan, Jean-Philippe F.
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                                Watanabe, Colin L. Wood, William I.
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ilarity 100.0%;
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CURRENT APPLICATION NUMBER: US/10/219,
CURRENT FILING DATE: 2002-08-12
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
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OR APPLICATION NUMBER: 60/123618
OR FILING DATE: 1999-03-10
IOR APPLICATION NUMBER: 60/125259
IOR APPLICATION NUMBER: 60/125775
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DR FILING DATE: 1998-09-24
DR FILING DATE: 1998-09-24
DR APPLICATION NUMBER: 60/101786
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DR FILING DATE: 1998-09-25
DR APPLICATION NUMBER: 60/101916
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DR FILING DATE: 1998-09-24
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R APPLICATION NUMBER: 60/10
R FILING DATE: 1998-09-16
RR APPLICATION NUMBER: 60/10
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R APPLICATION NUMBER: 60/10
R FILING DATE: 1998-10-28
R APPLICATION NUMBER: 60/10
R FILING DATE: 1998-10-29
RR APPLICATION NUMBER: 60/21
RR FILING DATE: 1998-10-30
R FILING DATE: 1998-10-30
OR FILING DATE: 1999-03-19

RAPPLICATION NUMBER: 60/125775

DR FILING DATE: 1999-03-23

OR FILING DATE: 1999-03-29

OR APPLICATION NUMBER: 60/127887

OR APPLICATION NUMBER: 60/127887

OR FILING DATE: 1999-04-05

OR APPLICATION NUMBER: 60/130232

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APPLICATION NUMBER: 60/1
FILING DATE: 1998-09-11
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FILING DATE: 1998-09-10
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FILING DATE: 1998-12-15
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FILING DATE: 1998-11-17
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FILING DATE: 1999-01-12
APPLICATION NUMBER: 60/119549
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FILING DATE: 1998-12-22
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FILING DATE: 1998-12-23
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OR APPLICATION NUMBER: 60/140723
OR FILING DATE: 1999-06-22
IOR APPLICATION NUMBER: 60/141037
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DR FILING DATE: 1999-12-07
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R APPLICATION NUMBER: 60/14
R FILING DATE: 1999-08-17
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R FILING DATE: 1999-08-17
R APPLICATION NUMBER: 60/15
R FILING DATE: 1999-08-31
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FILING DATE: 1999-05-14
APPLICATION NUMBER: 60/140650
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APPLICATION NUMBER: 60/
FILING DATE: 1999-04-27
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FILING DATE: 1999-07-28
APPLICATION NUMBER: 60/146963
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FILING DATE: 1999-07-20
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RESULT 33
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PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/079294
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PRIOR FILING DATE: 1998-03-25
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OR APPLICATION NUMBER: 60/090472
OR FILING DATE: 1998-06-24
OR APPLICATION NUMBER: 60/090557
OR FILING DATE: 1998-06-24
OR APPLICATION NUMBER: 60/090691
OR FILING DATE: 1998-06-25
OR APPLICATION NUMBER: 60/090695
OR FILING DATE: 1998-07-07
OR APPLICATION NUMBER: 60/091982
OR APPLICATION NUMBER: 60/095302
OR APPLICATION NUMBER: 60/095302
OR FILING DATE: 1998-08-04
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OR FILING DATE: 1998-06-17
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OR FILLING DATE: 1998-05-13
OR APPLICATION NUMBER: 60/085579
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OR FILING DATE: 1998-03-26
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OR APPLICATION NUMBER: 60/081819
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APPLICATION NUMBER: 60/082804
FILING DATE: 1998-04-22
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Gerritsen, Mary
Goddard, Audrey
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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OR APPLICATION NUMBER: 60/11558
OR APPLICATION NUMBER: 60/115565
OR FILING DATE: 1999-01-12
OR APPLICATION NUMBER: 60/115733
OR APPLICATION NUMBER: 60/115733
OR FILING DATE: 1999-01-12
OR FILING DATE: 1999-01-12 OR FILING DATE: 1998-09-18
OR APPLICATION NUMBER: 60/100919
OR FILING DATE: 1998-09-17
OR APPLICATION NUMBER: 60/101477
OR FILING DATE: 1998-09-23
OR APPLICATION NUMBER: 60/101738
OR FILING DATE: 1998-09-24
OR APPLICATION NUMBER: 60/101741
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OR APPLICATION NUMBER: 60/101786
OR FILING DATE: 1998-09-25
OR APPLICATION NUMBER: 60/101786
OR APPLICATION NUMBER: 60/101916 DR APPLICATION NUMBER: 60/106178
DR FILING DATE: 1998-10-28
DR APPLICATION NUMBER: 60/106248
DR APPLICATION NUMBER: 60/106464
DR APPLICATION NUMBER: 60/106905
DR APPLICATION NUMBER: 60/106905 OR APPLICATION NUMBER: 60/099803
OR FILING DATE: 1998-09-10
OR APPLICATION NUMBER: 60/099811
OR FILING DATE: 1998-09-10
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OR FILING DATE: 1998-09-10
OR APPLICATION NUMBER: 60/099816
OR FILING DATE: 1998-09-10
OR APPLICATION NUMBER: 60/10038
OR FILING DATE: 1998-09-11
OR APPLICATION NUMBER: 60/10038
OR FILING DATE: 1998-09-15
OR APPLICATION NUMBER: 60/10039
OR APPLICATION NUMBER: 60/10039
OR FILING DATE: 1998-09-15
OR APPLICATION NUMBER: 60/10039
OR FILING DATE: 1998-09-15
OR APPLICATION NUMBER: 60/10039
OR FILING DATE: 1998-09-15
OR APPLICATION NUMBER: 60/100627 AFILING DATE: 1998-11-18
APPLICATION NUMBER: 60/112422
FILING DATE: 1998-12-15
APPLICATION NUMBER: 60/113296
FILING DATE: 1998-12-22 R FILING DATE: 1998-08-11
R APPLICATION NUMBER: 60/0:
RFILING DATE: 1998-08-17
RR APPLICATION NUMBER: 60/0:
RR FILING DATE: 1998-08-26 FILING DATE: 1998-12-23 APPLICATION NUMBER: 60/113605 FILING DATE: 1998-11-17
APPLICATION NUMBER: 60/108849
FILING DATE: 170.00 FILING DATE: 1998-09-24 APPLICATION NUMBER: 60/ FILING DATE: 1998-09-24 FILLING DATE: 1998-11-03
APPLICATION NUMBER: 60/108787
FILING DATE: 1998-11-17
APPLICATION NUMBER: 60/108801 FILING DATE: 1998-09-16
APPLICATION NUMBER: 60/100848 APPLICATION NUMBER: 60/099596
FILING DATE: 1998-09-09
APPLICATION NUMBER: 60/099598
FILING DATE: 1998-09-09 APPLICATION NUMBER: 60/098544 FILING DATE: 1998-08-31 60/101922 60/097986 60/096791

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FILING DATE: 1999-04-27
APPLICATION NUMBER: 60/131445
FILING DATE: 1999-04-28
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FILING DATE: 1999-04-26
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FILING DATE: 1999-12-07
APPLICATION NUMBER: 60/169835
FILING DATE: 1999-12-07
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FILING DATE: 1999-11-16
APPLICATION NUMBER: 60/169445
FILING DATE: 1999-12-07
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AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
                                                              AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
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CURRENT FILING DATE: 2002-08-12
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR PILING DATE: 1997-09-17
PRIOR PILING DATE: 1997-09-17
PRIOR PILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
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PRIOR APPLICATION NUMBER: 60/063549
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SEQ ID NO 15
LENGTH: 1524
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PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079728
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PRIOR APPLICATION NUMBER: 60/
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                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Homo Sapien
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                   AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
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Grimaldi, J. Christopher
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Gerritsen, Mary
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                                                                                                                                                                                                 85.4%; Score 315; DE 100.0%; Pred. No. 2e-
Live 0; Mismatches
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; ORGANISM: Homo Sapien
US-10-219-062-15
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LENGTH: 1524
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/062287
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CURRENT FILING DATE: 2002-08-12
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170 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 229
                                                                                    30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
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OR ETLING DATE: 1997-10-28
OR APPLICATION NUMBER: 60/064103
OR FILING DATE: 1997-10-31
OR APPLICATION NUMBER: 60/069873
OR ETLING DATE: 1997-12-17
OR APPLICATION NUMBER: 60/078910
OR ETLING DATE: 1998-03-20
OR FILING DATE: 1998-03-20
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FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
FILING DATE: 1998-03-26
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                                                                                                                                               315;
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Smith, Victoria
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Gerritsen, Mary
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                                                                                                                                      100.0%; F1
                                                                                                                                                          85.4%; Score 315; 100.0%; Pred. No.
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LENGTH: 1524
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 246
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR EILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
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                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/079656 PRIOR FILING DATE: 1998-03-26
                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
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                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/079728
                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1998-03-25
                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/078910 PRIOR FILING DATE: 1998-03-20
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                                                                                                                                                                                           TYPE: DNA
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Local Similarity
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Grimaldi, J. Christopher
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Gerritsen, Mary
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                                                                              Conservative
                                                                                          100.0%;
                                                                     85.4%; Score 315; DB 42;
100.0%; Pred. No. 2e-159;
                                                                                                                                                                                                                                                                                                                                                                    60/079294
                                                                                                          Length 1524;
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PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR PPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
PRIOR PPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/069873
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR FILING DATE: 1998-03-20
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PRIOR FILING DATE: 1998-03-20
PRIOR FILING DATE: 1998-03-20
PRIOR FILING DATE: 1998-03-20
PRIOR FILING DATE: 1998-03-20
                                                                                                                     Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
     Matches 315; Conserv
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APPLICANT:
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                                                                                                                                                                                    PRIOR FILING DATE: 1998-03-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
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                                                                                      TYPE: DNA
ORGANISM: HOMO
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FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
                                                                                                                                                                                                                              FILING DATE: 1998-03-26
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Grimaldi, J. Christopher
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     Conservative
                  85.4%; Score 315; DB 42; 100.0%; Pred. No. 2e-159;
 0;
Mismatches
                                  DB 42;
 0;
                                Length 1524;
 Indels
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US-10-219-065-15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15; April 15
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PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063349
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PRIOR APPLICATION NUMBER: 60/064103
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                                                                                                                                   Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
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                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/079728 PRIOR FILING DATE: 1998-03-27
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TYPE: DNA
ORGANISM: Homo Sapien
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                                                                                                    LENGTH: 1524
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Grimaldi, J. Christopher
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Gerritsen, Mary
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                                                                            CURRENT APPLICATION NUMBER: US/10/219,066
CURRENT FILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR APPLICATION NUMBER: 60/079910
PRIOR APPLICATION NUMBER: 60/079910
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
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PRIOR APPLICATION NUMBER: 60/079294
PRIOR PILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079656
PRIOR APPLICATION NUMBER: 60/079656
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APPLICANT:
APPLICANT:
                                           PRIOR APPLICATION NUMBER: 60/079556
PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-27
PRIOR FILING DATE: 1998-03-27
                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3530P1C27
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                NUMBER OF SEQ ID
                                  Remaining Prior Application
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wood, William I.
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Gerritsen, Mary
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US-10-219-067-15
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US-10-219-066-15
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                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR PEPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
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                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND
TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                           PRIOR PRIOR
                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: P3530P1C51
CURRENT APPLICATION NUMBER: US/10/219,067
CURRENT FILING DATE: 2002-08-14
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nes 315; Conservative
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                                                                                                                                                         APPLICATION NUMBER: 60/063549
FILING DATE: 1997-10-28
APPLICATION NUMBER: 60/064103
FILING DATE: 1997-10-31
                                  FILING DATE: 1998-0: APPLICATION NUMBER:
                                                                                    APPLICATION NUMBER: 60/069873
FILING DATE: 1997-12-17
APPLICATION NUMBER: 60/078910
FILING DATE: 1998-03-20
                                                                     APPLICATION NUMBER: 60/079294
                    FILING DATE:
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Smith, Victoria
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Grimaldi, J. Christopher
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Gerritsen, Mary
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                     1998-03-26
                                                      1998-03-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Austin L.
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   60/079728
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; Remaining Prior Application data removed -
; NUMBER OF SEQ ID NOS: 246
; SEQ ID NO 15
; LENGTH: 1524
; TYPE: DNA
; ORGANISM: Homo Sapien
US-10-219-067-15
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                                   CURRENT APPLICATION NUMBER: US/10/219,068
CURRENT EILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-31
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1997-13-17
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: p3530P1C31
CURRENT APPLICATION NUMBER: US/10/219,068
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    APPLICATION NUMBER: FILING DATE: 1998-0
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Godowski, Paul J.
Grimaldi, J. Christopher
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Gerritsen, Mary
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NUMBER: 60/079294
1998-03-25
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Best Local Similarity
Matches 315; Conserv
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PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-27
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15, Applica GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: APPLICANT:
                                                                      TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC FILE REFERENCE: P3530p1C40

CURRENT APPLICATION UMMBER: US/10/219,069

CURRENT FILING DATE: 2002-08-13

PRIOR APPLICATION NUMBER: 10/119,480

PRIOR FILING DATE: 2002-04-09

PRIOR FILING DATE: 2002-09-13

PRIOR FILING DATE: 2002-09-17

PRIOR FILING DATE: 1997-09-17

PRIOR PILING DATE: 1997-09-17

PRIOR PPLICATION NUMBER: 60/062287

PRIOR PPLICATION NUMBER: 60/063549

PRIOR APPLICATION NUMBER: 60/063549

PRIOR FILING DATE: 1997-10-18

PRIOR APPLICATION NUMBER: 60/063549

PRIOR PILING DATE: 1997-10-28

PRIOR PILING DATE: 1997-10-28
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                                     APPLICATION NUMBER: 60/064103 FILING DATE: 1997-10-31
                 APPLICATION NUMBER: 60/069873
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Gerritsen, Mary
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APPLICANT: WOOD, WILLIAM I.

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND TITLE OF INVENTION: SECRETED AND THE SAME

FILE REFERENCE: P3530PIC42

CURRENT FILING DATE: 2002-08-13

PRIOR APPLICATION NUMBER: U5/10/219,070

PRIOR APPLICATION NUMBER: 60/059113

PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/062287

PRIOR APPLICATION NUMBER: 60/062287

PRIOR APPLICATION NUMBER: 60/063287

PRIOR APPLICATION NUMBER: 60/063549

PRIOR APPLICATION NUMBER: 60/063549

PRIOR FILING DATE: 1997-10-18
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APPLICANT: Godowski, Paul J.
APPLICANT: Gorimaldi, J. Christopher
APPLICANT: Gurney, Austin L.
APPLICANT: Smith, Victoria
APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079656
PRIOR APPLICATION NUMBER: 60/079656
PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-27
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Matches
                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Baker, K
APPLICANT: Desnoye
APPLICANT: Gerrits
APPLICANT: Goddard
APPLICANT: Godowsk
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SEQ ID NO 15
LENGTH: 1524
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les 315; Conserv
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Gerritsen, Mary
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2e-159;
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US-10-219-071-15
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US-10-219-070-15
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                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Baker, Kevin P.
APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Goddwski, Paul
APPLICANT: Grimaldi, J. Chi
APPLICANT: Grimaldi, J. Chi
APPLICANT: Grimaldi, J. Chi
APPLICANT: Grimaldi, J. Chi
APPLICANT: Smith, Victoria
APPLICANT: Stephan, Jean-Pi
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NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
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APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: 9353091C46
CURRENT APPLICATION NUMBER: US/10/219,071
CURRENT FILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
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PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
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Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Gerritsen, Mary
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100.0%; Pred. No. 2e:
Live 0; Mismatches
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US-10-219-072-15
Sequence 15, Application US/10219072
SEQUENCE 15, Application US/10219072
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Besnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Gerritsen, Mary
APPLICANT: Goddwski, Paul J.
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Gurney, Austin L.
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NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
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Best Local 9
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APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3530P1C45
CURRENT APPLICATION NUMBER: US/10/219,072
CURRENT FILING DATE: 2002-08-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/079728 PRIOR FILING DATE: 1998-03-27
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R FILING DATE: 1997-10-28

R FILING DATE: 1997-10-31

DR APPLICATION NUMBER: 60/069873

R FILING DATE: 1997-12-17

R FILING DATE: 1997-12-17
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APPLICATION NUMBER: 60/079294
FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
FILING DATE: 1998-03-26
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                                                                                               Goddard, Audrey
Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
Stephan, Jean-Philippe F.
Watanabe, Colin L.
Wood, William I.
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PRIOR FILING DATE: 1998-03-20
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PRIOR APPLICATION NUMBER: 60/079656
PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-27
PRIOR FILING DATE: 1998-03-27
Search completed: November Job time: 834.445 secs
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NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
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APPLICATION NUMBER: 60/062287
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US-10-240-425-140
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APPLICANT:

Godowski, Paul J. Grimaldi, J. Christopher

Watanabe, Colin L.

Stephan, Jean-Philippe Gurney, Austin L. Smith, Victoria

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RESULT 1 US-10-230-437-15 Sequence 15, Applica GENERAL INFORMATION: APPLICANT: Baker, K APPLICANT: Desnoye APPLICANT: Gerrits APPLICANT: Gerdard		44444033333333000000000000000000000000
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37-15 15, Application US/10230437 NFORMATION: T: Baker, Kevin P. T: Gesnoyers, Luc T: Gesnitsen, Mary T: Goddard, Andrev T: Goddard, Andrev		
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230437	ALIGNMENTS	US-60-417-190-48791 US-60-417-190-48792 US-60-417-190-48793 US-60-417-190-94485 US-60-417-190-94486 US-60-417-190-94486 US-60-417-190-94486 US-60-417-190-94489 US-60-417-190-94489 US-60-417-190-94489 US-09-513-999C-16171 US-09-513-999C-19578 US-09-513-999C-23316 US-09-513-999C-23316 US-09-513-999C-18526 US-09-513-999C-18527 US-09-513-999C-18527 US-09-513-999C-18527 US-09-513-999C-25794
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; TYPE: DNA
; ORGANISM: Homo
US-10-230-437-15
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PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-09-17
PRIOR PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/063349
PRIOR PILING DATE: 1997-10-28
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PRIOR PILING DATE: 1997-12-17
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PRIOR APPLICATION NUMBER: 60/078910
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PRIOR PILING DATE: 1998-03-20
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PRIOR APPLICATION NUMBER: 60/079656
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SEQ ID NO 15
LENGTH: 1524
          Query Match
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C94
CURRENT APPLICATION NUMBER: US/10/230,437
CURRENT FILING DATE: 2002-08-28
CURRENT FILING DATE: 10/110,490
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US-10-092-411A-2083/c

Sequence 2083, Application US/10092411A

GENERAL INFORMATION:

APPLICANT: Lynn Doucette-Stamm et al

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS

TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 032796-101

CURRENT APPLICATION NUMBER: US/10/092,411A

CURRENT FILING DATE: 1098-08-13

PRIOR APPLICATION NUMBER: US 09/134,001

PRIOR APPLICATION NUMBER: US 09/134,001

PRIOR TILING DATE: 1998-08-13

PRIOR FILING DATE: 1998-08-13
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; ORGANISM: Staphylococcus epidermidis
US-10-092-411A-1425
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CURRENT FILING DATE: 2002-03-07
PRIOR APPLICATION NUMBER: US 09/134,001
PRIOR FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5676
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Matches 315; Conser
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LENGTH: 1587
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PRIOR APPLICATION NUMBER: US 60/064,964
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; Mismatches 0;
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PRIOR EILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5676
SEQ ID NO 263
LENGTH: 393
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: Sequence 263, Application US/10092411A

: GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                             US-09-513-999C-36268/C
; Sequence 36268, Application US/09513999C
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-092-411A-263
                                                      ; TYPE: DNA; Homo say, ORGANISM: Homo say, US-09-513-999C-36268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 5
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Matches
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LENGTH: 156
TYPE: DNA
ORGANISM: Staphylococcus epidermidis
                                                                                                                      NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 36268
LENGTH: 111
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PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5676
       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Lynn Doucette-Stamm et al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCC TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 032796-101
CURRENT APPLICATION NUMBER: US/10/092,411A
CURRENT FILING DATE: 2002-03-07
                                                                                                                                                                                                                                                                                                                    APPLICANT: Duclert, A. APPLICANT: Giordano, J.Y. APPLICANT: Giordano, J.Y. TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins. FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Staphylococcus epidermidis
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18; Conserv
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            4.38;
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            Score 16;
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            Length 111;
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LENGTH: 397
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APPLICANT: Dumas Milne Edwards, J.B.
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SOFTWARE: Patent.pm
SEQ ID NO 28601
LENGTH: 433
TYPE: DNA
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TITLE OF INVENTION: Novel Murine Polynucleotide Tags and
TITLE OF INVENTION: Mutant Cells and Mutant Animals Observe Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing Street Processing 
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PRIOR FILING DATE: 1999-07-15
NUMBER OF SEQ ID NOS: 2908
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Zambrowicz, Bri
APPLICANT: Sands, Arthur T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24 PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Duclert, A. APPLICANT: Giordano, J.Y. TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins. FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature LOCATION: 275
                                                                                                                                                        OTHER INFORMATION: r-a or g
                                                                                                                                                                                                                       NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: n-a, g, c or t
                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
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                                     ME/KEY: misc_feature
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TION: r-a or g
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US-09-513-999C-3777/c
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; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                    US-09-513-999C-24773/c
; Sequence 24773, Application US/09513999C
; GENERAL INFORMATION:
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SOFTWARE: Patent.pm
SEQ ID NO 3777
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                                                                                                                                                             SOFTWARE: Patent.pm
SEQ ID NO 24773
                                                                                                                                                                             APPLICANT: Giordano, J.Y.

TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.

FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins. FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Duclert, A.
                                                                                                                                                                                                                                                                                                                             APPLICANT: Duclert, A. APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Dumas Milne Edwards, J.B.
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NAME/KEY: sig_peptide
LOCATION: 73..123
OTHER INFORMATION: score 9.3
OTHER INFORMATION: seq LALSLLILVLAFG/IP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
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LOCATION: 73..474
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                                                                                     TYPE: DNA ORGANISM: Homo sapiens FEATURE:
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                                                  NAME/KEY: misc_feature LOCATION: 182
NAME/KEY: misc_feature
                 FEATURE:
                                 OTHER INFORMATION: s=g or c
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                                                                                                                                            ENGTH: 250
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100.0%; Pred. No.
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US-09-513-999C-14500/c

; Sequence 14500, Application US/09513999C

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US-09-513-999C-1336
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; OTHER INFORMATION: k-g or t
US-09-513-999C-24773
                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                       Query Match
Best Local Similarity
                  APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Duclert, A.
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GENERAL INFORMATION:
     APPLICANT:
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TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
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PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
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LOCATION: 197
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LOCATION: 198
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LOCATION: 141..464
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Giordano, J.Y
               Duclert, A.
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100.0%; Pred. No.
ative 0; Mismatc
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Pred. No.
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US-10-240-425-140
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LENGTH: 579
                                                                            Query Match
Best Local (
                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/240,425
CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: PCT/US01/09847
PRIOR FILING DATE: 2001-03-28
PRIOR APPLICATION NUMBER: US 60/193,446
PRIOR FILING DATE: 2000-03-31
NUMBER OF 5EQ ID NOS: 1588
SOFTWARE: PATENTIN VET: 2.1
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SEQ ID NO 14500
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TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5026
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APPLICANT: Boland, Joseph
APPLICANT: Lord, Reginald
APPLICANT: Alvarez, Chris
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                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
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LOCATION: 325
OTHER INFORMATION: r=a or
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OTHER INFORMATION: y=c
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                             304 GCCAAGGCCCAAGAA 318
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81 GCCAAGGCCCAAGAA 95
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Local Similarity 100.08;
es 15; Conservation
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Scherf, Uwe
                                                              Conservative
                                                                          4.1%; Score 15;
100.0%; Pred. No.
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Pred. No.
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33;
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; LCCATION: (1)..(686)
; OTHER INFORMATION: n - a or c or g
US-10-240-425-710
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PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 1588
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 710
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TITLE OF INVENTION: Gene Expression Profiles in Esophageal Tissue
FILE REFERENCE: 44921-5026
CURRENT APPLICATION NUMBER: US/10/240,425
CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: PCT/US01/09847
PRIOR FILING DATE: 2001-03-28
PRIOR FILING DATE: 2001-03-28
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Scherf, Uwe
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Lord, Reginald V.
Alvarez, Chris
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CHAWLA, Narinder K.
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                                                                                                                                                                                                                                                                                                       YANG,
                                                              LEE, Ernestine A.
LEE, Soo Yeun
ELLIOTT, Vicki S.
                                                                                                                                                          GRIFFIN, Jennifer A.
                                                                                                                                                                                                                                         ELLIOTT, Vicki S.
               GORVAD, Ann E
KABLE, Amy E.
                                                                                                              BAUGHN, Mariah R.
                                                                                                                                                                           EMERLING,
                                              RAMKUMAR, Jayalaxmi
                                                                                                                             ZEBARJADIAN,
                                                                                                                                            LAL, Preeti G.
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                                                                                                                                                                                                                                                                                      Yuming
Dyung Aina M
                                                                                                                                                                                                                                                                                                                                                                                                 Henry
                                                                                                                                                                                        Monique G.
                                                                                             Ernestine A.
                                                                                                                                                                                                                                                                                                       Junming
                                                                                                                                                                                                                                                                                                                                                                                    Bridget A.
                                                                                                                                                                                                                                                                                                                                     Brendan M.
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100.0%;
                                                                                                                                                                           Brooke M.
                                                                                                                                                                                                                                                                                                                                                                     Patricia M.
                                                                                                                             Yeganeh
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Pred. No.
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PRIOR FILING DATE: 2001-10-03
PRIOR APPLICATION NUMBER: US 60/343,718
PRIOR FILING DATE: 2001-10-19
PRIOR APPLICATION NUMBER: US 60/343,980
PRIOR FILING DATE: 2001-11-02
PRIOR APPLICATION NUMBER: US 60/332,426
PRIOR FILING DATE: 2001-11-16
NUMBER OF SEQ ID NOS: 64
                                                                         RESULT 16
US-10-264-237-1045
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Best Local Similarity
Matches 15; Conserv
                                    Sequence 1045, Application US/10264237 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 3.1
SEQ ID NO 561
LENGTH: 1292
                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                        Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: PCT/US02/32032
CURRENT FILING DATE: 2002-10-03
APPLICANT: Birse et
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Nuc
FILE REFERENCE: PAl31P1
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NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: 7340485CB1
                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens
FEATURE:
REATURE:
NAME/KEY: misc_feature
LOCATION: (534)..(534)
OTHER INFORMATION: n equals a,t,g, or c
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                                                                                                                                                                                                                                    Local Similarity
es 15; Conserv
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100.08; P1
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  Nucleic Acids, Proteins, and Antibodies
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Pred. No.
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US-10-125-923A-555/c; Sequence 555, Application US/10125923A; GENERAL INFORMATION:
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                                                                                                         RESULT 18
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GENERAL INFORMATION:
APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA131p1
CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
                                                                                                                                                                                                                                   Matches
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SEQ ID NO 1045
LENGTH: 1542
                  APPLICANT: Baker, Kevin P. APPLICANT: Chen, Jian
   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
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PRIOR TILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
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SOFTWARE: PatentIn Ver. 3.
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CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (1459)..(1459)
OTHER INFORMATION: n equals a,t,g, or c
                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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LOCATION: (1471)...(1471)
OTHER INFORMATION: n equals a,t,g, or
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Desnoyers, Luc
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100.0%; Pred. No.
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Pred. No.
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; ORGANISM: Homo sapiens US-10-264-237-1252
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                                                                                                                           SOFTWARE: PatentIn Ver. 3.1
SEQ ID NO 1252
LENGTH: 1913
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1252, Application:
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LENGTH: 1773
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CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/205,515 PRIOR FILING DATE: 2000-05-19
                                                                                                                                                                                                                                                                                                                                       APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA131P1
                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 2876
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PRIOR APPLICATION NUMBER: 60/063120
PRIOR FILING DATE: 1997-10-24
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PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059266
PRIOR FILING DATE: 1997-09-18
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                                                                                                         TYPE: DNA
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4.18;
100.08;
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Pred. No.
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Pred.
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                     Length 1913;
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RESULT 20
US-10-240-851-69/c
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PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 09/543,771
PRIOR ETLING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 09/229,319
PRIOR FILING DATE: 1999-01-13
PRIOR APPLICATION NUMBER: US 60/071,449
PRIOR APPLICATION NUMBER: US 60/071,449
PRIOR APPLICATION NUMBER: US 60/105,511
PRIOR APPLICATION NUMBER: US 60/105,511
PRIOR APPLICATION NUMBER: US 60/105,511
PRIOR FILING DATE: 1998-10-23
RESULT 22
US-10-240-425-1461
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                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1, Application US/10192280
GENERAL INFORMATION:
APPLICANT: Dattagupta, Nanibhushan
APPLICANT: Shah, Ketan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DETECTING
TITLE OF INVENTION: STREPTOCOCCUS AGALACTIAE CAPSULAR POLYSACCHARIDE SYNTHESIS
TITLE OF INVENTION: GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 69, Application US/10240851 GENERAL INFORMATION:
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SEQ ID NO 69
LENGTH: 3513
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SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 25020
TYPE: DNA
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Best Local Similarity 100.0%;
Matches 15; Conservative
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/240,851 CURRENT FILING DATE: 2002-10-04
                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: GENES
FILE REFERENCE: 475412001300
CURRENT APPLICATION NUMBER: US/10/192,280
CURRENT FILING DATE: 2002-07-09
                                                                          10941 AAGAAGTGATGGAGC 10955
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Local Similarity 100.0%;
nes 15; Conservation
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Alvarez, Chris

APPLICANT: Wetzel, Jon C.

APPLICANT: Vockley, Joseph G.

TITLE OF INVENTION: Gene Expression Profiles in Esophageal Tissue

FILLE REFERENCE: 44921-5026

CURRENT APPLICATION NUMBER: US/10/240,425

CURRENT APPLICATION NUMBER: US/10/240,425

CURRENT FILING DATE: 2001-03-28

PRIOR FILING DATE: 2001-03-28

PRIOR FILING DATE: 2001-03-28

PRIOR FILING DATE: 2001-03-31

NUMBER OF SEQ ID NOS: 1588

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 1461

TYPE: DNA

ORGANIC**
                                                                                                                                                                                                                                     APPLICANT: Boland, Joseph F.
APPLICANT: Lord, Reginald V.
APPLICANT: Lord, Reginald V.
APPLICANT: Wetzel, Jon C.
APPLICANT: Wetzel, Jon C.
APPLICANT: Wetzel, Jon C.
APPLICANT: Vockley, Joseph G.
APPLICANT: Vockley, Joseph G.
TITLE OF INVENTION: Gene Expression Profiles in Esophageal Tissue
FILE REFERENCE: 44921-5026
CURRENT APPLICATION NUMBER: US/10/240,425
CURRENT APPLICATION NUMBER: US/2001-03-09-30
PRIOR APPLICATION NUMBER: PCT/US01/09847
PRIOR APPLICATION NUMBER: US 60/193,446
PRIOR APPLICATION NUMBER: US 60/193,446
PRIOR APPLICATION NUMBER: US 60/193,446
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 1588
SOFTWARE: PatentIN Ver. 2.1
LENGTH: 74822
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                                                                                                                                                ; OTHER INFORMATION: Genbank Accession No. AL049798 US-10-240-425-1121
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Best Local Similarity
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Best Local
                                                                       Matches
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                                                                                                                                                                                           TYPE: DNA ORGANISM: Homo sapiens
70056 ATGTGTCAGAAAGAA 70070
                                                                                                                                                                                     FEATURE:
                   137 ATGTGTCAGAAAGAA 151
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                                                                       Local Similarity 100 les 15; Conservative
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100.0%; Pred. No.
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Pred. No.
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39;
                                                                                          DB 6; Length 74822; 40;
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Sequence 48788, Application US/60417190
GENERAL INFORMATION:
APPLICANT: Giulia Kennedy
APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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US-60-417-190-48787/c
; Sequence 48787, Appl
; GENERAL INFORMATION:
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US-60-417-190-48789/c
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US-60-417-190-48788
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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEO ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEO ID NO 4878
LENGTH: 25
TYPE: DNA
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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V
SEQ ID NO 48787
       SEQ ID NO 48789
LENGTH: 25
                                                                                                                                                                                                                      Sequence 48789, Application US/60417190 GENERAL INFORMATION:
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Best Local 9
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                                            CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
                                                                                                                        APPLICANT: Giulia Kennedy
APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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Local Similarity 100.0%;
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Pred. No.
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Pred. No.
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US-60-417-190-48792/c
; Sequence 48792, Application US/60417190
; GENERAL INFORMATION:
; APPLICANT: Giulia Kennedy
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                                                                        RESULT 29
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US-60-417-190-48790/c
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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 48791
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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 48790
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Best Local
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Matches 14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Giulia Kennedy
APPLICANT: Hajime Matsuzaki
APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Wethods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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Pred. No.
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Pred. No.
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US-60-417-190-48793/c ; Sequence 48793, Application US/60417190 ; GENERAL INFORMATION:
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APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 48792
LENGTH: 25
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                                                                                                                                   US-60-417-190-94484
                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 94484
LENGTH: 25
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CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 48793
LENGTH: 25
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APPLICANT: Hajime Matsuzaki
APPLICANT: Hei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
Matches
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APPLICANT: Hajime Matsuzaki
APPLICANT: Hajime Matsuzaki
APPLICANT: Hei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
TILE REFERENCE: 3522
CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
                                                                                                                                                            TYPE: DNA ORGANISM: Homo sapien
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100.0%; Pred. No.
                               3.8%; Score 14;
100.0%; Pred. No.
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Sequence 94487, Application US/60417190

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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 94485
LENGTH: 25
TYPE: DNA
ORGANISM: Homo sapien
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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 94486
LENGTH: 25
TYPE: DNA
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GENERAL INFORMATION:
APPLICANT: Giulia Kennedy
APPLICANT: Hajime Matsuzaki
APPLICANT: Hei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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GENERAL INFORMATION:
APPLICANT: Giulia Kennedy
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APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms
FILE REFERENCE: 3522
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Local Similarity 100.0%; Pred. No.
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                                                                                                                                 ; ORGANISM: Homo sapien US-60-417-190-94489
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                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 94489
LENGTH: 25
                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                Sequence 94489, Application US/60417190 GENERAL INFORMATION:
                                                                                                Query Match
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                                                                                                                                                                                                                                                                               APPLICANT: Giulia Kennedy
APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 94487
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APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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                                                                                                                                                                LENGTH: 25
TYPE: DNA
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TYPE: DNA
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TYPE: DNA
                               143 CAGAAAGAAGTGAT 156
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19 CAGAAAGAAGTGAT 6
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                                                                 14;
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100.0%; Pred. No.
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                                                                             Score 14; DB 7; Length 25; Pred. No. 1e+02;
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. 1e+02;
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: Sequence 19578, Application US/09513999C
: GENERAL INFORMATION:
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US-09-513-999C-19578
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US-09-513-999C-17785, Application US/09513999C
; Sequence 17785, Application US/09513999C
; GENERAL INFORMATION:
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US-09-513-999C-16171/c
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SEQ ID NO 17785
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APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Glordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
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Best Local (
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SEQ ID NO 16171
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CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
CURRENT FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins
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APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
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100.0%; Pred. No. 1.1
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; ORGANISM: Homo sapiens
US-09-513-999C-19578
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                          US-09-513-999C-18596
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NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 19578
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SEQ ID NO 23316
LENGTH: 112
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SEQ ID NO 18596
LENGTH: 116
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Best Local
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Query Match
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CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 199-02-26
NUMBER OF SEQ. ID NOS: 36681
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                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24 PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26
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APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins
FILE REFERENCE: 59.USZ.REG
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                                                                                                                                              NUMBER OF SEQ ID NOS: 36681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
                                                 TYPE: DNA
ORGANISM: Homo
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Pred. No.
   Score
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   14;
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   DВ
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PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 14544
PLENCTH: 125
TYPE: DNA
ORGANISM: Homo sapiens
US-09-513-999C-14544
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US-09-513-999C-14544/c
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Sequence 18527, Application US/09513999C
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
RESULT 44
US-09-513-999C-28218
; Sequence 28218, Application US/09513999C
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
                                                                                                                                                                                                                                                                                                                    ; LENGTH: 129
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-513-999C-18527
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SOFTWARE: Patent.pm
SEQ ID NO 18527
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Best Local Similarity
Matches 14; Conserv
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Best Local Similarity 100.0%;
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TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Dumas Milne Edwards, J.B.
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100.0%; Pred. No. 1.1e+02;
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Pred. No. 1.1e+
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Page 12

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APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
CURRENT FILING DATE: 2000-02-24
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 25794
LENGTH: 142
TYPE: DNA
ORGANISM: Homo sapiens
US-09-513-999C-25794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Duclert, A.
APPLICANT: Glordano, J.Y.
APPLICANT: Glordano, J.Y.
FILE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 28218
LENGTH: 138
LENGTH: 138
CORGANISM: Homo sapiens
US-09-513-999C-28218
Search completed: November 8, 2002, 02:02:17 Job time: 69.3091 secs
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US-09-513-999C-25794
Sequence 25794, Application US/09513999C
GENERAL INFORMATION:
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OM nucleic - nucleic search, using sw model
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Score Match Length DB ID  315 85.4 704 13 BI919074 315 85.4 707 13 BI913989 315 85.4 920 9 AL538562 314 85.1 593 10 BE386060 307 83.2 747 13 BI917149
h Length DB ID  4 704 13 B1919074  4 707 13 B1913989  4 920 9 AL538562  1 593 10 BE386060  2 747 13 B1917149  9 987 14 BQ666420

FEATURES

Location/Qualifiers

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## ALIGNMENTS

cDNA Library Preparation: Life Technologies, Inc.	Tissue Procurement: Life Technologies, Inc.	Email: cgapbs-r@mail.nih.gov	EST.	BI919074.1 GI:16200128		mRNA sequence.	603180881F1 NIH_MGC_121 Homo sapiens cDNA clone IMAGE:5244956 5',	BI919074 704 bp mRNA linear EST 16-OCT-2001
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLAM11618 row: a column: 21 High quality sequence start: 4	CDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLAM11618 row: a column: 21 High quality sequence start: 4	Tissue Procurement: Life Technologies, Inc. CDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLAM1618 row: a column: 21 High quality sequence start: 4	Homo sapiens  Homo sapiens  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 704)  NIH-MGC http://mgc.nci.nih.gov/.  National Institutes of Health, Mammalian Gene Collection (MGC)  Unpublished (1999)  Contact: Robert Strausberg, Ph.D.  Email: cgapbs-r@mail.nih.gov  Tissue Procurement: Life Technologies, Inc.  CDNA Library Preparation: Life Technologies, Inc.  CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)  DNA Sequencing by: Incyte Genomics, Inc.  Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov  Plate: LLAM1618 row: a column: 21  High quality sequence start: 4	human.  Homo sapiens  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 704)  NIH-MGC http://mgc.nci.nih.gov/.  National Institutes of Health, Mammalian Gene Collection (MGC)  Unpublished (1999)  Contact: Robert Strausberg, Ph.D.  Email: ggapbs-r@mail.nih.gov  Tissue Procurement: Life Technologies, Inc.  CDNA Library Preparation: Life Technologies, Inc.  CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)  DNA Sequencing by: Incyte Genomics, Inc.  Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:  http://image.llnl.gov  Plate: LLAM11618 row: a column: 21  High quality sequence start: 4	BIG19074.1 GI:16200128  EST. human. Homo Sapiens Homo Sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Metazoa; Chordata; Craniata; Hominidae; Homo.  1 (bases 1 to 704) NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999) Contact: Robert Strausberg, Ph.D. Email: cgapbs-rémail.nih.gov Tissue Procurement: Life Technologies, Inc. cDNA Library Preparation: Life Technologies, Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLAM1618 row: a column: 21 High quality sequence start: 4	iens a; Metazoa; Eutheria; Eutheria; I to 704) http://mgc. Institutes hed (199) Robert Str gapbs-remant rocurement; brary Prepa brary Array uencing by: istribution rough the I LAMN1618 In. 19 Lity sequer	mRNA sequence. B1919074 B1919074 B1919074.1 GI:16200128 EST. human. Homo Sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 704) NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999) Contact: Robert Strausberg, Ph.D. Email: gapbs-r@mail.nih.gov Tissue Procurement: Life Technologies, Inc. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. 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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
                                                                                                                                     Contact: Robert Strausberg, Ph.D.
                                                                                                                                                      NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 707)
                                                                                                                                                                                                                                                                                                                                                                BI913989 707 bp mRNA linear EST 16-OCT-20 603180565F1 NIH_MGC_121 Homo sapiens cDNA clone IMAGE:5244569 5',
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/db_xref="taxon:9606"
/clone="IMAGE:5244956"
/clone_lib="NIH_MGC_121"
/lab_host="DH10B"
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100.0%;
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Pred. No. 1.7e-166;
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                                                                                                                                     AUTHORS
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435 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 494
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                                                                                                                                                                   Full-length cDNA libraries Unpublished (2001)
                                                                                                                                  Contact: Genoscope
                                                   Email: segref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
                                                                                 BP 191 91006 EVRY cedex - France
                                                                                                          Genoscope - Centre National de Sequencage
                                                                                                                                                                                                                             Li,W.B., Gruber,C., Jessee,J. and Polayes,D
                                                                                                                                                                                                                                                                                    Mammalia; Eutheria; Primates; Catarrhini;
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                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Location/Qualifiers
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Location/Qualifiers
1. .920
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/clone="IMAGE:5244569"
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601276753F1 NIH_MGC_20 Homo sapiens
                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                 Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                    NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
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                              cDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note-"Organ: Fetal brain; Vector: pCMVSPORT 6; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the
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week, 24 week and 26 week)"
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100.0%; Pred. No. 1.8e-166;
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.A.G.E. Consortium/LLNL at: image.llnl.gov
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                       BI917149 747 bp mRNA linear EST 16-OCT-2001 603181571F1 NIH_MGC_121 Homo sapiens cDNA clone IMAGE:5245648 5',
                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies,
                                                                                                                                                                  NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                  Homo sapiens
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                              BI917149.1 GI:16181111
                                                                                                                                                                                                                                                                                                                                                                   mRNA sequence.
BI917149
                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                   Unpublished (1999)
               cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                           (bases 1 to 747)
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/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: skin; Vector: pTB7; Site_1: XhoI; Site_2:
cloned into EcoRI/XhoI sites using the following 5'
adaptor: GGCACGAG(G)... Size-selected >500bp for average
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/clone_lib="NIH_MGC_20"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     307;
                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                           NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC:
                                                    Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                      Mammalia; Eutheria; Primates; Catarrhini; Losses 1 to 987)
                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                               BQ668420
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Plate: LLAM11619 row: n column: 17
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/clone="IMAGE:5245648"
/clone_lib="NIH_MGC_121"
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Preparation: Rubin Laboratory
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Pred. No. 5.7e-162;
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3C_102 Homo !
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ne IMAGE:6275121
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TITLE
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                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                  Unpublished (1999)
                                                                                                                                                                                                                                NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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http://image.llnl.gov
Plate: LLCM2457 row: m column: 10
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DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
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/clone_lib="NIH_MCC_102"
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/tissue_type="epidermoid carcinoma, cell line"
/lab_host="NHIDB (phage-resistant)"
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/db_xref="taxon:9606"
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Pred. No. 3.
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                                                                                                                                           AUTHORS
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Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM11792 row: p column: 10
High quality sequence stop: 690.
Location/Qualifiers
                                                              Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Bukaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo.

1 (bases 1 to 340)

1 (bases 1 to 340)

1 (bases 1 to 340)

M., Lolark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trewaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R.
                                                                                                                                                                                                                                                                                                H06756 340 bp mRNA linear EST 21-y183b07.rl Soares infant brain INIB Homo sapiens cDNA clone IMAGE:44619 5', mRNA sequence.
 Washington University Scho 4444 Forest Park Parkway,
                                  Unpublished (1995)
Contact: Wilson RK
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/tissue_type="hypothalamus"
/lab_host="DH10B"
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/db_xref="taxon:9606"
/clone="IMAGE:5312889"
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Pred. No. 5.5e-149;
School of Medicine way, Box 8501, St.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: est@watson.wustl.edu
Insert Size: 1737
High quality sequence stops: 312
Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 1737
Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence stop: 312. Location/Qualifiers
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                                                                                                                                                                             1010 bp 602018318F1 NCI_CGAP_Brn67 Homo 5', mRNA secuence
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 1010)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                        Homo sapiens
                                                                                                                                            BF345141.1 GI:11292606
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/db_xref="GDB:417160"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CATCAGCTGCTACCACCCCTCTTTGTAAC 301
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                271;
                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                         Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                               NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                     mRNA sequence.
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Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM9422 row: d column: 05
High quality sequence stop: 649.
Location/Qualifiers
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Tissue Procurement: David N. Louis, M.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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cDNA Library Preparation: Ling Hong/Rubin Laboratory
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/note="Organ: brain; Vector: pCMV-SPORT6; Site_1:
Site_2: Sall; Cloned unidirectionally. Primer: Oli
Average insert size 2.3 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
a 344 c 254 g 146 t
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/clone="IMAGE:4153684"
/clone_lib="NCI_CGAP_Brn67"
/tissue_type="anaplastic oligodendroglioma with 1p/19q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
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1 (bases 1 to 948)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                   Contact: Robert Strausberg, Email: cgapbs-r@mail.nih.gov Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                       Unpublished (1999)
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AGENCOURT_8627686 NIH_MGC_43
CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
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Clone distribution: MGC clone distribution information can be
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Plate: LLCM2492 row: k column:
Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
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National Institutes of Health, Mammalian
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               Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image_llnl.gov
                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 570)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                mRNA sequence.
BI831919
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603078792F1 NIH_MGC_119 Homo sapiens cDNA clone IMAGE:5170606 5',
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                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
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Plate: LLAM11424
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/clone="IMAGE:5181585"
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Pred. No. 2.7e-134;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      237;
                             For clone availability, additional sequence and expression information related to this EST, please check the TIGR Human Gene Index (http://www.tigr.org/tdb/hgi/hgi.html) Seq primer: M13 Reverse.
                                                                                                   Email: arkerlav@tigr.org
                                                                                                                                                        The Institute for Genomic Research 9712 Medical Center Drive, Rockvil
                                                                                                                                                                                                    Bioinformatics
                                                                                                                                                                                                                   Contact: Kerlavage, AR
                                                                                                                                                                                                                                     Other_ESTs: EST56848 THC106611
                                                                                                                                                                                                                                                                                                    Adams,M.D., Kerlavage,A.R., Fields,C. and Venter,J.C. 3,400 expressed sequence tags identify diversity of transcripts
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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           Location/Qualifiers
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/clone_lib="NIH_MGC_119"
/tissue_type="medulla"
/lab_host="DH10B"
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                                                                                                                                                                                                                                                                     found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLCM1759 row: e column: 01 High quality sequence stop: 585.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1999)
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, N
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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            /Clone_TIMAGE:4876392"
/Clone_lib="NIH_MGC_43"
/Clone_lib="NIH_MGC_43"
/tissue_type="normal pigmented retinal epithelium"
/tissue_type="normal pigmented retinal epithelium"
/lab host="DH10B (phage-resistant)"
/note="Organ: eye; Vector: pOTB7; Site_1: xhoI; Site_2: EcoRI; cDNA made by oligo-dT priming Directionally cloned into EcoRI/XhoI sites using the following 5 adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
                                                                                                                                                                                                 /db_xref="taxon:9606"
                                                                                                                                                                                                                  /organism="Homo sapiens"
                                                                                                                                                                                                                                                         Location/Qualifiers
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/db_xref="ARCC (Inhost):150796"
/db_xref="taxon:9606"
/clone_lib="Infant brain"
/sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="Organ: brain; Vector: lafmid BA; Site_1: HindIII;
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
Plate: LLCM/60 row: 1 column: 21
High quality sequence stop: 651.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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198 c 165 g 163 t
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/db_xref="taxon:9606"
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Pred. No. 4.3e-115;
      Score 205;
Pred. No.
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                   209 TGTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTT 268
                                                                        427 GAAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCC
                                                                                                              149 GAAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCC 208
                                                                                                                                                                     Local
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TGTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis, M.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MCC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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BI554034
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 698)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Plate: LLAM11784 row: d column: 08
                                                                                                                                                                     Similarity
                                                                                                                                                 53.1%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         quality sequence stop:
                                                                                                                                                                                                                                            /lab_host="DH10B (T1 phage-resistant)"
/note="Organ: brain: Vector: pCMV-SPORT6; Site_1
Site_2: Sali; Cloned unidirectionally. Primer: C
Average insert size 2.3 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
a 231 c 204 g 132 t
                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:9606"
/clone="IMAGE:5309527"
/clone_11b="NCI_CGAP_Brn67"
                                                                                                                                                                                                                                                                                                                                                                         tissue_type="anaplastic oligodendroglioma with 1p/19q/
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Pred. No. 2.8e-99;
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imer: Oligo dT
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AW954549
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                                                                                                                                                                    AGCTGCTGCAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCC 334
                                                                                                                                                TCGGCCCTCA 344
                                                                                                                       TCGGCCCTCA 245
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                mRNA sequence.
AA582576
                                       nn55e03.sl NCI_CGAP_Kid6 Homo sapiens
     AA582576.1
                                                        AA582576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Assessment of gene expression patterns in a model of colon tumor metastasis using a 19,200 element cDNA microarray Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: John Quackenbush
The Institute for Genomic Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hegde, P., Q1, R., Abernathy, K., Dharap, S., Gaspard, R., Ga, I.E., Saeed, A.I., Sharov, V., Lee, N.H., Yeatman, T.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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Mammalia; Eutheria;
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301 838 3528
301 838 0208
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                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="MAGE resequences, MAGC"
/note="Vector: pBluescriptSKm"
/note="134 g 134 t
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     GI:2359936
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                                                     mRNA
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                                       NA linear EST 26-SEP-1997 cDNA clone IMAGE:1087804 3',
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Tumor Gene Index
                                                                          Homo sapiens
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                                                                                                                                         mRNA sequence.
                                                                                                                                                                   AI767459
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290 CCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCCCTCA 344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCCCTCA 432
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.National Cancer Institute, Cancer Genome Anat
                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 522)
                                                                                                                                                                                                                                                                                        wh26h03.x1 NCI_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:2381909
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cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-GGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 904 Std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 407.
                                                                                                                                                                                                                     AI767459.1 GI:5233968
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Tissue Procurement: L. Jeffrey Medeiros, M.D.,
Emmert-Buck, M.D., Ph.D.
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/db_xref="taxon:9606"
/clone="IMAGE:1087804"
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100.0%; Pred. No. 1.8e-87;
^. wismatches 0;
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Genome Anatomy Project (CGAP)
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MAGE:2381909 3',
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCCCTCA 460
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Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
                                                                             Genome Re
97044477
                                                                                                                                                                                                                                                                                                                               BM974798 541 bp mRNA linear EST 21-M
UI-CF-EC1-acd-i-10-0-UI.sl UI-CF-EC1 Homo sapiens cDNA clone
UI-CF-EC1-acd-i-10-0-UI 3', mRNA sequence.
                                                        Contact: McCray, PB
                                                                                                                    discovery
                                                                                                                                    Normalization and subtraction: two approaches
                                                                                                                                                          Bonaldo, M.F.,
                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 541)
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                                      McCray Lab
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                      University of Iowa
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University of Iowa
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                                                                                               Res. 6 (9), 791-806 (1996)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone="IMAGE: 2381909"
/clone_lib="NCI_CGAP_Kid11"
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                                                                                                                                                          Lennon, G. and Soares, M.B.
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  Med Labs, Iowa City, IA 52242, USA
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FEATURES
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Tissue Procurement: Dr. M. J. Welsh, University of Iowa
CDNA Library preparation: Dr. M. Bento Soares, University of Iow
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                 AV717094
AV717094
AV717094
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Fax: 319 356 7171
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1 (bases 1 to 651)
Xu, X., Gu, J., Liu, F
Song, H., Gu, Y., Yan
                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                             Homo sapiens
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/note="Organ: Lung: Vector: pT73-Pac (Pharmacia) with a
modified polylinker; Site_1: EcoR I; Site_2: Not I;
UT-CF-ECI is a normalized cDNA library containing the
following tissue(s): Normal lung from adult and from fetal
day 64, day 87, week 19 and week 42. The library was
constructed according to Bonaldo, Lennon and Soares,
Genome Research, 6:791-806, 1996 First strand cDNA
synthesis was primed with an oligo-dT primer containing a
Not I site. Double stranded cDNA was ligated to an EcoR I
adaptor, digested with Not I, and cloned directionally
into pT73-Pac vector. The oligonucleotide used to prime
the synthesis of first-strand cDNA contains a library tag
sequence that is located between the Not I site and the
                                                                                                                                                                                                                         DCB Homo sapiens
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/db_xref="taxon:9606"
/clone="UI-CF-EC1-acd-i-10-0-UI"
/clone_lib="UI-CF-EC1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TAG_SEQ=AAGTGCTTAC"
1 142 c 120 g
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/dev_stage="Adult and Fetal"
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Liu,F., Qu,J., Zhao,M., Li,Y., Hua'., Yang,Y., Gao,G., Xiao,H., Li,N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      47.4%; Score 175; DB 14; 100.0%; Pred. No. 1.9e-87;
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DCBAZD04 5', mRNA sequence.
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       Qian, B.,
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                            Zhou,J.
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         Gao, X.,
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        Email: paul-mccray@uiowa.edu
Tissue Procurement: Dr. M. J. Welsh, University of Iowa
cDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                                                                                                                                                                                                                                    BM980701 652 bp mRNA linear EST 21-M/UI-CF-EN1-add-j-04-0-UI.S1 UI-CF-EN1 Homo sapiens cDNA clone UI-CF-EN1-add-j-04-0-UI 3', mRNA sequence.
                                                                                                                                                                                              Contact: McCray, PB
                                                                                                                                                                                                                              Genome Res. 6 (9),
                                                                                                                                                                                                                                               discovery
                                                                                                                                                                                                                                                                             1 (bases 1 to 652)
Bonaldo, M.F., Lenno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong,
T01: 86-21-50801919(ex.45)
                                                                                                                                                                 University of Iowa
                                                                                                                                                                                   McCray Lab
                                                                                                                                                                                                                                                          Normalization and subtraction: two approaches
                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: hanzg@chgc.sh.cn
This clone is available at CHGC in Shanghai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fax: 86-21-50801922
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Homo sapiens cDNA DCB clones
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                                                                                                               University of Iowa Med Labs, Iowa City, IA 52242, USA 319 356 4866
319 356 7171
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(www.resgen.com).
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/lab_host="BM25.8"
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100.0%; Pred. No. 2e-87;
Live. 0; Mismatches
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178 g    146 t
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                                                                                          JOURNAL
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                         University of Iowa
                                         McCray Lab
                                                     Contact: McCray, PB
                                                                                                          discovery
                                                                           97044477
                                                                                          Genome
                                                                                                                                         Bonaldo, M.F.,
                                                                                                                                                                                                      Homo sapiens
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412 CCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCA 466
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                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 738)
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UI-CF-EN1-adh-b-10-0-UI.S1 UI-CF-EN1 Homo sapiens
UI-CF-EN1-adh-b-10-0-UI 3', mRNA sequence.
                                                                                                                                                                                                                     Normalization and subtraction: two approaches
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/note="Organ: Lung; Vector: pT7T3-Pac (Pharmacia) with a
modified polylinker: Site_1: EcoR I; Site_2: Not I;
UI-CF-ENI is a normalized cDNA library containing the
following tissue(s): Primary Lung Cystic Fibrosis
Epithelial Cells. The library was constructed according to
Bonaldo, Lennon and Soares, Genome Research, 6:791-806,
1996. First strand cDNA synthesis was primed with an
oligo-dT primer containing a Not I site. Double stranded
cDNA was ligated to an EcoR I adaptor, digested with Not
I, and cloned directionally into pT7T3-Pac vector. The
oligonucleotide used to prime the synthesis of
first-strand cDNA contains a library tag sequence that is
located between the Not I site and the (dT)8 tall. The
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TAG_LIB-UI-CF-EN1
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/clone_lib="UI-CF-EN1"
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100.0%; Pr
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RESULT 25
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Best Local Similarity
                                                                                                                                           ORGANISM
                                        AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 175;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutherla; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 890)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                   601763551F1 NIH_MGC_20 Homo sapiens
                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                   BF125425.1 GI:10964465
                                                                                                                                                                                                                                                              mRNA sequence.
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Tissue Procurement: Dr. M. J. Welsh, University of Iowa
cDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Researchers may obtain clones from Research
Genetics (www.resgen.com).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note-*Organ: Lung; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: ECOR I; Site_2: Not I; UI-CF-EN1 is a normalized cDNA library formation the following tissue(s): Primary Lung Cystic Fibrosis Epithelial Cells. The library was constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806,
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/clone_lib="UI-CF-EN1"
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/db_xref="taxon:9606"
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Pred. No. 2.1e-87;
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one IMAGE:4026252 5',
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KEYWORDS
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               qe22c09.x1 Soares_fetal_lung_NbHL19W Homo sapiens cDNA clone IMAGE:1739728 3', mRNA sequence.
                                                                                                                                                                            Unpublished (1997)
                                                                                                                                                                                                Tumor Gene Index
                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 449)
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                                                                                                                                                                                                              NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1999)
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/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: skin: Yector: pOTB7; Site_1: XhoI; Site_2:
ECORI; CDNA made by oligo-dT priming. Directionally
cloned into ECORI/XhoI sites using the following 5;
adaptor: GGCACCANG(6). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
32 a 258 c 239 g 161 t
Location/Qualifiers
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/clone="IMAGE:4026252"
/clone_lib="NIH_MGC_20"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      47.2%; Score 174; DB 12; 100.0%; Pred. No. 8.2e-87;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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170 GGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGG 229
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACC 289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGG 336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACC 396
                                                                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutherla; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 878)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BF125664.1 GI:10964704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           601763237F1 NIH_MGC_20 Homo sapiens
                                                                                                                                          High quality sequence stop: 598.
Location/Qualifiers
                                                                                                                                                                                              CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                             Unpublished (1999)
                                                                                                                                                                             Plate: LLCM856 row: i column:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="taxon:9728"
/clone_lib="Soares_fetal_lung_NbHL19W"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /dev_stage="19 weeks"
/lab_host="DH10B (ampicillin resistant)"
/tissue_type="melanotic melanoma"
/lab_host="DH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
                                                                      /organism="Homo sapiens"
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/clone="IMAGE:4026320"
                                                    /clone_lib="NIH_MGC_20"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             146 AAAGAAGTGATGGAGCAAAGTGCCGGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCG 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       206 GCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGG 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          360 AAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCG 419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: est@watson.wustl.edu
Insert Size: 1628
High quality sequence stops: 326 Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 1628
Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                   Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1995)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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   /db_xref="GDB:405062"
/db_xref="taxon:9606"
/clone="IMAGE:32715"
                                                                                                                              /clone_lib="Soares infant brain lNIB
                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                               /sex="female'
                                                                                                                                                                                                   /organism="Homo sapiens"
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100.0%; Pred. No.
100.0%; Pred. No.
100.0%; Pred. No.
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Pred. No. 7.3e-83;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hawkins, M., Holman
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r EST 17-APR-1995 cDNA clone

Waterston

REFERENCE AUTHORS

TITLE

JOURNAL

SOURCE KEYWORDS VERSION ACCESSION DEFINITION

EST

BF125664 mRNA sequence.

ORGANISM

Homo sapiens

COMMENT

FEATURES

source

ø Not Not RESULT 27 BF125664

BF125664

õ В ş 망 Š

290

230

277

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397

ECORI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)." a 194 c 251 g 155 t

Length 878; Indels 0, Gaps

86 GACTGCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAAACGTTCAAGACATGTGTCAG 145

300 GACTGCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAG 359

BASE COUNT ORIGIN

101

Query Match Best Local Similarity

Matches

173;

Conservative

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                                                                                                                                                                                                                                                                                                                                          Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1810
Fax: 314 286 1810
Email: est@watson.wustl.edu
Insert Size: 1633
High quality sequence stops: 287
Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 1633
Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R. The Washu-Merck EST Project
                                                                                                                                                                                                                                                                                                           High quality sequence stop: 287.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpubl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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1 (bases 1 to 429)
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429 bp mRNA linear EST 29-, ym37f07.rl Soares infant brain lNIB Homo sapiens cDNA clone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ished (1995)
            double-stranded cDNA was ligated to Hind III adaptors (Pharmacla), digested with Not I and directionally cloned into the Not I and Hind III sites of the Lafmid BA vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo." a 126 c 103 g 92 t 1 others
      double-stranded
                                                                                                                        /dev_stage="73 days post natal"
                                                                                                                                                                                     /db_xref="taxon:9606"
/clone="IMAGE:50205"
                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="GDB:423015"
                                                                                                                                                 /sex="female"
                                                                                                                                                              /clone_lib="Soares infant brain lNIB"
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                                                                                                                                                                                                                                                                                      ocation/Qualifiers
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Pred. No. 2.1e-82;
cDNA was ligated to Hind III adaptors
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                           cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consorttum/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 787 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mRNA seq
AI299246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tissue Procurement: Christopher Moskaluk, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tumor Gene Index
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              α
                        /clone="IMAGE:1900098"
/clone_lib="NCI_CGAP_Kid5"
/tissue_type="2 pooled tumors (clear cell type)"
/lab_host="DHIOB"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Pharmacia), digested with Not I and directionally cloned into the Not I and Hind III sites of the Lafinid BA vector. Library went through one round of normalization. Library constructed by Bento Soares and M. Fatima Bonaldo."

127 c 108 g 107 t 5 others
                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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Pred. No. 2.2e-82;
Pred. No. 2.2e-82;
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99 g
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sapiens cDNA clone IMAGE:1900098 3',
             127 t
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 960 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 420.
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qd98e07.x1 Soares_testis_NHT Homo sapiens cDNA clone
3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NGI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Anat
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1 (bases 1 to 424)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1737540"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="Soares_testis_NHT"
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100.0%; Pred. No. 6.6e-74;
                                                                          100.0%;
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                                                   0
                                                Score 148; DB 9;
Pred. No. 3.2e-72;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs r@mail.nih.gov

The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail cDNA Library Preparation: M.B. Soares Lab Clone distribution: NCI-CGAP clone distribution information can be found through the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AWU148U2
UI-H-BIO-aag-b-04-0-UI.s1 NCI_CGAP_Sub1 Homo sapiens cDNA clone IMAGE:2709078 3', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
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National Cancer Institute, Cancer Genome Ana
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           / ARD_HOST DATASE (MARGE CINNOLOGYES)
/ ARD_HOST DATASE (MARGE CIONELDS 12391, 1420-1431)
/ ARD_HOST DATASE (MARGE CIONELDS 13, 2391, 1475-3, 2871-281)
/ ARD_HOST DATASE (MARGE CIONELDS 13, 2871-281)
/ ARD CONSTITUTES A MIXTURE of 21 normalized or subtracted NCI_CGAP_INTER (MCI_CGAP_CO16, NCI_CGAP_PC22, NCI_CGAP_PC38, NCI_CGAP_CO16, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_CO16, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_CO16, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, NCI_CGAP_EC4, N
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/db_xref="taxon:9606"
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/clone_lib="NCI_CGAP_Subl"
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NCI_CGAP_Co10 pool 1 LLAM 2644-2653, 2871-2872
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REFERENCE
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KEYWORDS
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AA776783
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                        www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 941 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 510.
                                                                                                                                                                                                                                                                                                                                                                                  CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1997)
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National Cancer Institute, Cancer Genome Anat
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 516)
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EST.
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subtracted library contained 530,000 recombinants.
Subtraction was performed as previously described [Bonaldo, Lennon & Soares (1996): Normalization and Subtraction:
Two Approaches To Facilitate Gene Discovery. Genome
Research 6,791-806.
TAG_LIB-NCI_CGAP_Kid5
                                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="1276662"
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                                                                                                                                                                        /sex="male"
                                                                                                                                                                                           /clone_lib="Soares_testis_NHT"
                                                                                                                                                                                                                                                                                        Location/Qualifiers
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Pred. No. 1.6e-70;
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TITLE
                       Query Match
Best Local
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al Similarity
143; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                        cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov Plate: LLCM543 row: h column: 17 High quality sequence stop: 642.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (1999)
Contact: Robert St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                         /tissue_type="choriocarcinoma"
/lab_host="DH10B (phage-resistant)"
/note="Organ: placenta; vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Size-selected >500bp
for average insert size 1.8kb. Library constructed by
Ling Hong in the laboratory of Gerald M. Rubin (University
of California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
a 227 c 217 g 155 t
                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:9606"
/clone="IMAGE:3845440"
/clone_lib="NIH_MGC_21"
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139 c 121 g 139 t
                                                                                                                                                                                                                                                                                                                                                                                  organism="Homo sapiens"
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                  38.8%;
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              Score 143; DB 12;
Pred. No. 2.6e-69;
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Pred. No. 2.3e-69;
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                                   Length 794;
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Best Local
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                                                                                                                   AAGTGATGGAGCAAAGTGCCGGG 172
                      AAGTGATGGAGCAAAGTGCCGGG 426
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BE730800.1 GI:10144792
EST.
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601570755F1 NIH_MGC_21 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                             Score 143; DB 12;
Pred. No. 2.8e-69;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 78
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Creation of genome-wide protein expression libraries using random activation of gene expression
activation of gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3201 Carnegie Ave, Cleveland,
Tel: 216 431 9900
Fax: 216 361 9596
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RST33311 Athersys RAGE Library
BG213596
BG213596.1 GI:13735283
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                                                                                                                              86184021 484 bp mRNA linear EST 21-APR-
RST2936 Athersys RAGE Library Homo sapiens cDNA, mRNA sequence.
BG184021
BG184021.1 GI:13705708
EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Scott J. Cain
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                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                               Homo sapiens
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  (bases 1 to 484)
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/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression
/note="See 'Creation of Genome-wide Protein Expression',
Libraries using Random Activation of Gene Expression',
Nature Biotechnology, in press. Note that even though the
cell type indicated is HT1080, since a random activation
method was used, these sequence tags are not necessarily
method was used, these sequence tags are not necessarily
expressed in HT1080 under normal circumstances."
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Pred. No. 2.8e-68;
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                                                                                                                                                             Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 781)
Harrington,J.J., Sherf,B., Rundlett,S., Jackson,P.D., Perry,R., Cain,S., Leventhal,C., Thornton,M., Ramachandran,R., Whittington,J., Lerner,L., Costanzo,D., McElligott,K., Boozer,S., Mays,R., Smith,E., Veloso,N., Kilka,A., Hess,J., Cothren,K., Lo,K., Offenbacher,J., Danzig,J. and Ducar,M.
                                                                                                                                                                                                                                                                                                                                                                                                                        BG193459
RST12592 Athersys RAGE Library
BG193459
                                                                                                                Creation of genome-wide protein expression libraries using random activation of gene expression
                                        Athersys,
                                                          Contact: Scott J. Cain
                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Harrington,J.J., Sherf,B., Rundlett,S., Jackson,P.D., Perry,R., Cain,S., Leventhal,C., Thornton,M., Ramachandran,R., Whittington,J., Lerner,L., Costanzo,D., McElligott,K., Boozer,S., Mays,R., Smith,E., Veloso,N., Klika,A., Hess,J., Cothren,K., Lo,K., Offenbacher,J., Danzig,J. and Ducar,M.
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Nat. Biotechnol. 19 (5), 440-445 (2001)
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Creation of
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Location/Qualifiers
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Tel: 216 431 9900
Fax: 216 361 9596
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Scott J. Cain
Carnegie Ave, Cleveland, OH 44115, USA 216 431 9900
                                                                                                    Biotechnol. 19 (5),
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/db_xref="taxon:9606"
/db_xref="taxon:9606"
/clone_lib="Athersys RAGE Library"
/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression
/note="See 'Creation of Genome-wide Protein Expression
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                                                                                                  440-445 (2001)
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1 (bases 1 to 424)

Harrington,J.J., Sherf,B., Rundlett,S., Jackson,P.D., Perry,R., Cain,S., Leventhal,C., Thornton,M., Ramachandran,R., Whittington,J., Lerner,L., Costanzo,D., McElligott,K., Boozer,S., Mays,R., Smith,E., Veloso,N., Klika,A., Hess,J., Cothren,K., Lo,K., Offenbacher,J., Danzig,J. and Ducar,M.

Creation of genome-wide protein expression libraries using random
                                                                                                                                                                                                                                                                                                                                                 3201 Carnegie Ave, Cleveland, OH 44115, USA
Tel: 216 431 9900
Fax: 216 361 9596
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Scott J. Cain
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BG188218.1 GI:13709905
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High quality sequence stop: 322
Location/Qualifiers
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/db_xref="taxon:9606"
/clone_lib="Athersys RAGE Library"
/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression /, note="See vising Random Activation of Gene Expression', Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation
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/cell_line="HT1080"
/note="see 'Creation of Genome-wide Protein Expression
/Inote="see 'Creation of Genome-wide Protein Expression',
Ilbraries using Random Activation of Gene Expression',
Nature Biotechnology, in press. Note that even though the
cell type indicated is HT1080, since a random activation
method was used, these sequence tags are not necessarily
expressed in HT1080 under normal circumstances."
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/db_xref="taxon:9606"
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Best Local Similarity
Matches 140; Conserv
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Athersys, Inc.
Athersys, Inc.
Athersys, Inc.
Athersys, Inc.
Athersys, Inc.
Athersys, Inc.
Tel: 216 431 9900
Fax: 216 361 9596
Email: scain@athersys.com
High quality sequence stop: 323.
High quality sequence stop: 323.
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Harrington,J.J., Sherf,B., Rundlett,S., Jackson,P.D., Perry,R., Cain,S., Leventhal,C., Thornton,M., Ramachandran,R., Whittington,J., Lerner,L., Costanzo,D., McElligott,K., Boozer,S., Mays,R., Smith,E., Veloso,N., Klika,A., Hess,J., Cothren,K., Lo,K., Offenbacher,J., Danzig,J. and Ducar,M.
Creation of genome-wide protein expression libraries using random activation of gene expression
Nat. Biotechnol. 19 (5), 440-445 (2001)
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RST12050 Athersys RAGE Library Homo sapiens CDNA, mRNA sequence.
BG192926
BG192926.1 GI:13714613
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Scott J. Cain
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                                                                                                                                                                  /note-"see 'Creation of Genome-wide Protein Expression Libraries using Random Activation of Gene Expression', Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation method was used, these sequence tags are not necessarily expressed in HT1080 under normal circumstances."
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/db_xref="taxon:9606"
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100.0%; Pred. No. -
100.0%; Pred. No. -
11ve 0; Mismatches
                                                                                           37.7%;
                                                                          Score 139; DB 12;
s; Pred. No. 3.6e-67;
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                                                                                                             Length 323;
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Qy     34       Db     1       Qy     94       Db     61       Qy     154       Db     121	Query Ma Best Loc Matches	BASE COUNT	FEATURES SOUTCE	JOURNAL MEDLINE COMMENT	TITLE	REFERENCE AUTHORS	RESULT 41 BG210457 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM	Qy 154 c Db 121 c	Oy 94 C
CTTTGCGCTGCAAATCCAGTGCTACCAGTGTAAAGAATTCCAGCTGAACAACGACTGCTC 93	Match 37.7%; Score 139; DB 12; Length 323; Local Similarity 100.0%; Pred. No. 3.6e-67; Local Similarity 100.0%; Pred. No. 3.6e-67; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	/organism="Homo sapiens" /db_xref="taxon:9606" /clone_lib="Athersys RAGE Library" /cell_line="HT1080" /note="see 'Creation of Genome-wide Protein Expression /note="see 'Creation of Genome-wide Protein Expression', Libraries using Random Activation of Gene Expression', Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation method was used, these sequence tags are not necessarily expressed in HT1080 under normal circumstances." 90 a 83 c 78 g 72 t	3201 Carnegie Ave, Cleveland, OH 44115, USA Tel: 216 431 9900 Fax: 216 361 9596 Email: scain@athersys.com High quality sequence stop: 323. Location/Qualifiers e 1. 323	9 (5), 440-445 (2001) Cain	E. Veloso, N. Kilka, A. Hess, J., Cothren, K., Lo, K., Offenbacher, J., Danzig, J. and Ducar, M. Creation of genome-wide protein expression libraries using random activation of gene expression.	Mammalia; Eutheria; Primates; Ctatarrhini; Hominidae; Homo.  1 (bases 1 to 323)  1 (bases 1, Sherf, B., Rundlett, S., Jackson, P.D., Perry, R.,  Harrington, J.J., Sherf, B., Rundlett, S., Jackson, R.D., Perry, R.,  Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J.  Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J.  Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J.  Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J.  Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J.  Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J.  Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J.	BG210457 RST30002 Athersys RAGE Library Homo sapiens cDNA, mR BG210457 BG210457.1 GI:13732144 EST. human. Homo sapiens	GATGGAGCAAAGTGCCGGG 172 	CTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTCAAGACATGTGTCACAGAAAGATCT [

RESULT 42 BG215210 LOCUS DEFINITION

BG215210 324 bp mRNA linear EST 21-APR-2 RST34874 Athersys RAGE Library Homo sapiens cDNA, mRNA sequence.

EST 21-APR-2001

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BG200048/c
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, u.,
                 Harrington, J.J., Sherf, B., Rundlett, S., Jackson, P.D., Perry, R., Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, Lerner, L., Costenzo, D., WcElligott, K., Boozer, S., Mays, R., Smit, E., Veloso, N., Klika, A., Hess, J., Cothren, K., Lo, K., Offenbacher
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Creation of genome-wide protein expression libraries using random activation of gene expression
Nat. Bistechnol. 19 (5), 440-445 (2001)
                                                                                                                                                                                                                                                               BG200048 330 bp mRNA linear EST 21-APR-7
RST19350 Athersys RAGE Library Homo sapiens cDNA, mRNA sequence.
                                                                                                                                                                       Homo sapiens
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                                                                                                                                               Eukaryota; Metazoa;
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Fax: 216 361 9596
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/note="See 'Creation of Genome-wide Protein Expression
/Inter-see 'Creation of Genome-wide Protein Expression',
Libraries using Random Activation of Gene Expression',
Nature Biotechnology, in press. Note that even though the
cell type indicated is HT1080, since a random activation
method was used, these sequence tags are not necessarily
expressed in HT1080 under normal circumstances."
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/db_xref="taxon:9606"
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Primates;
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                                                 Athersys, Inc.
3201 Carnegie Ave, (
Tel: 216 431 9900
Fax: 216 361 9596
                                                                                                                                                       Harrington, J.J., Sherf, B., Rundlett, S., Jackson, P.D., Perry, R., Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J., Lerner, L., Costanzo, D., McElligott, K., Boozer, S., Mays, R., Smith, E., Veloso, N., Klika, A., Hess, J., Cothren, K., Lo, K., Offenbacher, J., Danzig, J. and Ducar, M.

Creation of genome-wide protein expression libraries using random activation of gene expression
Nat. Biotechnol. 19 (5), 440-445 (2001)
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RST815 At
BG181955
Email: scain@athersys.com
High quality sequence stop: 331.
Location/Qualifiers
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                                                                                                                                Contact: Scott J. Cain
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Tel: 216 431 9900
Fax: 216 361 9596
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Nat. Biotechnol. 19 (5), 440-445 (2001)
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Athersys RAGE Library Homo
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/db_xref="taxon:9606"
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Primates;
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Pred. No. 3.7e-67;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Creation of genome-wide protein expression libraries using random activation of gene expression Nat. Biotechnol. 19 (5), 440-445 (2001)
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BG204666.1 GI:13726337
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Location/Qualifiers
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1 (bases 1 to 331)
                                                                                                                                                                                                                                                                                                                                 Email: scain@athersys.com
                                                                                                                                                                                                                                                                                                                                                        Tel: 216 431 9900
Fax: 216 361 9596
                                                                                                                                                                                                                                                                                                                                                                                                    3201 Carnegie Ave, Cleveland, OH 44115, USA
                                                                                                                                                                                                                                                                                                                                                                                                                         Athersys, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Scott J. Cain
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             Libraries using Random Activation of Gene Expression', Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation method was used, these sequence tags are not necessarily expressed in HT1080 under normal circumstances."

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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

ORGANISM REFERENCE AUTHORS TITLE JOURNAL	RESULT 1 AX093381 LOCUS DEFINITION ACCESSION VERSION KEYWORDS		C 44 45	43	c 41	c 40		C 36		3 2 2		c 29	C 28		0 0 224			c 19	18		c 14			10	ი დ დ	70	νΩ	ى 4	, 10 (	<u> </u>	Result	
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                        Sequencing, Maryland;
Gaithersburg, Maryland;
Web site: http://www.nisc.nih.gov/
                                                               cDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: National Institutes of Health Intramural
Sequencing Center (NISC),
Shevchenko,Y., Wetherby,K.D., Beckstrom-Sternberg,S.M.,
                                                                                                                                                                                                                                  Submitted (05-NOV-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer
                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                    NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
                                                                                                                                                                                                                                                                          Direct Submission
                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Homo sapiens, clone MGC:29643 IMAGE:3641660, mRNA, complete
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KEYWORDS VERSION

AX358762.1 GI:18675282

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RESULT 3
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Sequence 15 from Patent WO0193983 AX358762
                                             AX358762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Clone distribution: MGC clone distribution information can be through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.g. Series: IRAL Plate: 39 Row: b Column: 19
This clone was selected for full length sequencing because it passed the following selection criteria: Hexamer frequency ORR analysis, Similarity but not identity to protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Benjamin,B., Blakesley,R.W., Bouffard,G.G., Brinkley,C., Brooks,S., Dietrich,N.L., Guan,X., Gupta,J., Ho,S.-L., Karlins,E., Legaspi,R., Lim,M., Maduro,Q.L., Masiello,C., Mastrian,S.D., McCloskey,J.C., McDowell,J., Pearson,R., Snyder,B., Stantripop,S., Thomas,P.J., Tiongson,E.E., Touchman,J.W., Tsurgeon,C., Vogt,J.L., Walker,M.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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292 c 261 g 196 t
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/protein_id="AAH17318.1"
/protein_id="AAH17318.1"
/db_xref="Gi:16678240"
/translation="MWVLGIAATFCGLFLLPGFALQIGCYQCEEFQLNNDCSSPEFIV
/translation="MWVLGIAATFCGLFLLPGFALQIGCYQGEEFGLNNDCSSPEFIV
/TVNNDTNYQDMCQKEVMBQSAGIMYRKSGASSAACLLASAGYQSFCSPGKLNSVCISCCN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="Pancreas,
/clone_lib="NIH_MGC_39"
/lab_host="DH10B-R"
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/db_xref="taxon:9606"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent: WO 0193983-A 15 13-DEC-2001;
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                                                                                                   Baker, K.P., Desnoyers, L., Gerritsen, M.E., Goddard, A., Goddowski, P.J., Grimaldi, J.C., Gurney, A.L., Smith, V., Watanabe, C.K. and Wood, W.I.
                                                                                                                                                                                                                                                                                       Sequence 15 from Patent Ax362255
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                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                 Secreted and transmembrane polypeptides and nucleic
                               Genentech,
                                              Patent: WO 0208288-A 15 31-JAN-2002;
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                                                                                                                          Smith, V., Stephan, J.P.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                           AK094501.1 GI:21753575 oligo capping; fis (full insert sequence). Homo sapiens alzheimer cortex cDNA to mRNA, clone_lib:BRALZ2 clone:BRALZ2001350.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AK094501 1832 bp mRNA linear PRI 15-JUL-2002
Homo sapiens cDNA FLJ37182 fis, clone BRALZ2001350, weakly similar
to Homo sapiens Gz-selective GTPase-activating protein (RGS20)
Isogai,T. and Yamamoto,J.

Direct Submission

Direct Submission

Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7

Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7

Kazusa-Kamatari, Kisarazu, Chiba 292-0812, Japan

Kazusa-Kamatari, Kisarazu, Chiba 292-0812, Japan

Kezusa-Kamatari, Kisarazu, Chiba 192-0812, Japan

Construction: Helix Research Institute (HRI) (Supported by Japan

Construction: Helix Research Institute (HRI) (Supported by Japan

Key Technology Center etc.); 5'- & 3'-end one pass sequencing: RAB,
                                                                                                                                                                                                                                       Kawakami,B., Sugiyama,A., Takemoto,M., Sugiyama,T., Irie,R., Otsuki,T., Sato,H., Wakamatsu,A., Ishii,S., Yamamoto,J., Isono Kawai-Hio,Y., Saito,K., Nishikawa,T., Kimura,K., Yamashita,H., Matsuo,K., Nakamura,Y., Sekine,M., Kikuchi,H., Kanda,K., Wagatsuma,M., Murakawa,K., Kanehori,K., Takahashi-Fujii,A., Oshima,A., Suzuki,Y., Sugano,S., Nagahari,K., Masuho,Y., Nagai
                                                                                                                                                                                                      Unpublished
                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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433 c 435 g
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Pred. No. 1.1e-95
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Masuho,Y., Nagai,K.

Isono, Y.,

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SOURCE
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                                                                               AUTHORS
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                                                                                                                                                                                                   1890 bp
Sequence 203 from Patent EP1067182.
AX136281
              Patent: EP 1067182-A 203 10-JAN-2001;
Helix Research Institute (JP)
                                                             Ota,T., Isogai,T., Nishikawa,T., Kawai,Y.,
Hayashi,K.
                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                            Secretory protein or membrane protein
                                                                                                                                       Homo sapiens
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                                                                                            (bases 1 to 1890)
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Location/Qualifiers
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/db_xref="G1:21753576"
/db_xref="G1:21753576"
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TPLCNGPRPKKRGSSASALRPGLRTTILFLKLALFSAHC"
501 c 513 g 407 t
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/clone_lib="BRALZ2"
/note="cloning vector: pME18SFL3"
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/db_xref="taxon:9606"
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                                                                       Sugiyama,T.
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                                                                                                                Ovarian tumor sequences and methods of Patent: WO 0118046-A 9 15-MAR-2001; CORIXA CORPORATION (US)
                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1. (bases 1 to 396)
                                                                                                                                                                 Xu, J. and Stolk, J.A.
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                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
1. .396
        /note="n = A,T,C or G"
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TPLCNQPRPKKRGSSASALRPGLRTTILFLKLASSRHTAELKEMPPPPALFFQPSPPT
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274. .771
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Best Local :
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Best Local Similarity
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Sequence 3 from Patent W00181634.
AX319944
                                                                                                                                                                                                                                                                                                                                                                                                                                            Galvin, K.A. and Rudolph-Owen, L.A. Methods and compositions for the diagnosis and treatment of cardiovascular and tumorigenic disease using 4941 patent: WO 0181634-A 3 01-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AX319944.1 GI:17901491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                             Millennium Pharmaceuticals, Inc. (US)
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                                                                                                                            /Translation="maspslpgsdcsqlidhshuperevatmiktitlivyklifykg
/translation="maspslpgsdcsqlidhshuperevitimplits
sytlsgkihtplepacsyatilhyltlsferviaicheprykavsgecoykiligetw
vtsalvalpilfapacsyatilhyltlsferviaicheprykavsgecoykiligetw
vtsalvalpilfapacsyatilhyltlsferviaicheprykavsgecoykiligetw
vtsalvalpilfapacsyatilhyltlypshercunhscythheopetsumsictnissrwf
vtossitegarvylvylvbapacwnnmydvlmksgkgslaggtrppoliksgesetsty
arrotiffaltivyllavcwnenoirrimaaakpkhowfrsytraymillefsetfty
lssvinpllytyssoofrryfydylccrlslohahekelryndakstydsaravcher
                                                                                                                                                                                                                                                         /protein_id="CAD19382.1"
/db_xref="GI:17901492"
                                                                                                                                                                                                                                                                                                                                                                      /organism-"Homo sapiens"
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Pred. No. 2.8e-89;
                                                                        677 g
   Score 336.4;
Pred. No. 3.
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                                                                      Submitted (13-APR-2000) Katsuyuki Hashimoto, National Institute of Infectious Diseases, Division of Genetic Resources; 23-1, Toyama 1-chome, Shinjuku-ku, Tokyo 162-8640, Japan (E-mail:khashi@nih.go.jp, URL:http://www.nih.go.jp/yoken/genebank/, Tel:81-3-5285-1111(ex.2120), Fax:81-3-5285-1181)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fis (full insert sequence).
Mus musculus (strain:C57BL) adult female cDNA to mRNA,
clone_lib:Sugano mouse brain mncb clone:MNCb-0671.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 isolation of full-length cDNA clones from mouse brain
                                                                                                                                                                                                                                                                        Lab host:
                                                            part of this sequence is reported
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                                                                                                                                                                                                                                                                                         Name:
                                                                                                                                                                                                                                                                                                   http://www.nih.go.jp/yoken/genebank/
/organism="Mus musculus"
/strain="C57BL"
                                           location/Qualifiers
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ORIGIN
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AX136556
             BASE COUNT
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ORGANISM
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ORIGIN
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                                                                                                                                                  Ota, T., Isogai, T., Nishikawa, T., Kawai, Y.,
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                         /db_xref="taxon:9606"
                                         organism="Homo sapiens"
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NCTVNVQDMCQKEVTEQSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRFKKRGSSASAIRFBCLLTTLLFFHLALCLAHC"
a 487 c 511 g 408 t
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/dev_stage="adult"
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/db_xref="GI:7670500"
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                                                                                    Direct Submission

Submitted (09-JAN-2002) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 6 On Aug 9, 2001 this sequence version replaced gi:14488388.
                                                                                                                                                                                                                                                    Direct Submission
Submitted (09-AUG-2001) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St. Louis,
                                                                                                                                                                                                                       MO 63108, USA
5 (bases 1 to 129676)
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Submitted (10-SEP-2000) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St.
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               Web site: http://genome.wustl.edu/gsc
Contact: sapiens@watson.wustl.edu
------ Summary Statistics
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The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoegawa,K., Woon,P.Y., Zhao,B., Frengen,E., Tateno,M., Catanese,J.J. and de Jong,P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at the Roswell Park Cancer Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >- 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The clone sequenced to the left is RP11-1L22; the clone sequenced to the right is RP11-159N20, 2000 bp overlap. Actual start of thi clone is at base position 1 of RP11-258B17; actual end is at base position 60003 of RP11-159N20.
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Location/Qualifiers
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                                                  /rpt_family="(GA)n"
32667. .32787
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27926. .2
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26656. .2704
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                                                                                     32053
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8. .21796
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AC124493/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      166 TGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGC 225
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Sequencing vector: M13; 08
Sequencing vector: plasmid; 1008
Chemistry: Dye-primer ET; 08 of reads
Chemistry: Dye-terminator Big Dye; 1008 of reads
Assembly program: Phrap; version 0.990319
Consensus quality: 208484 bases at least Q40
                                                                                                                                                                                                                                                                                                                                                                Submitted (05-JUL-2002) Genome Sequencing Center, 4444 Forest Park
Parkway, St. Louis, MO 63108, USA
On Jul 5, 2002 this sequence version replaced gi:21426614.
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McPherson, J.D. and Waterston, R.H.
Direct Submission
                                                                                                                                      Summary Statistics
                                                                                                                                                                                      Center project name: M_BA0462P13
                                                                                                                                                                                                                                 Web site:http://genome.wustl.edu/gsc/index.shtmlContact: submissions@watson.wustl.edu
                                                                                                                                                                                                                                                                              Center: Washington University Genome Sequencing Center Center code: WUGSC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished
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McPherson, J.D. and Waterston, R.H.
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                                                                                                                                                                                                              ------ Project Information ------
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35361. .35676
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                                                                                                                                                                                                                                                   TGGCTCCGCACCACCATCCT 369
                                                       Homo sapiens BAC clone RP11-159N20
AC010974
AC010974.9 GI:19033964
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Homo sapiens
                   Homo sapiens.
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* This record will be updated with the finished sequence
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49555. 107384
/note="assembly_name:Contig11"
107485. 209885
/note="assembly_name:Contig12"
44784 c 46164 g 58745 t
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28718. .49454
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11432. .28617
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/chromosome="UNK"
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11431: gap of unknown length
28617: contig of 17186 bp in length
28717: gap of unknown length
49454: contig of 20737 bp in length
49554: gap of unknown length
107384: contig of 57830 bp in length
107484: gap of unknown length
107484: gap of unknown length
209885: contig of 102401 bp in length.
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Pred. No. 2e-35;
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5 (bases 1 to 147131)
Waterston, R.
Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VanBrunt,A. and Strowmatt,C.
The sequence of Homo sapiens BAC clone RP11-159N20
Unpublished (2001)
3 (bases 1 to 147131)
NEIGHBORING SEQUENCE INFORMATION:
The clone sequenced to the right is RP11-458A7, 200 bp overlap; clone sequenced to the right is RP11-258B17. Actual start of the clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of RP11-159N20; actual end is at base position 1 of
                                                                                                                                                                                                                                                                                                                              Research Genetics, Inc. (http://www.resgen.com) or Pieter de and coworkers at http://www.chori.org
VECTOR: pBACe3.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOURCE INFORMATION:

The RPCI-11 human BAC library was made from the blood of one donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Freng Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 5::1-8. The clone may be obtained either
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
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8227. .8260
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /rpt_family="MIR"
118. .1013
             /note="match to EST 15183. .15321
                                                                          /note="match to EST 15183. .15321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="2"
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                                             /note="match to EST BE386060
15183. 15321
                                                                                                                                       /rpt_family="MER1_type"
13456. .13776
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12505
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                             BG704350 (NID:g13977603)"
                                                                                       AI239596 (NID:g3834993) qh37b10.x1"
                                                                                                                                                                                                                                                  BG187166 (NID:g13708853)"
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RESULT 14
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                                 Rattus norvegicus clone CH230-140B23, ***, 69 unordered pieces.
              AC110334
AC110334.3 GI:21738206
HTGS_PHASE1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="similar to Homo sapiens EST BE963800
(NID:g11767217)"
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yj75b04.s1"
18938. .19409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="match to EST BG184021 (NID:g13705708)"
15184. .15320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="similar to Bos taurus EST BE750540 (NID:g10164532)"
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15184. .15321
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16665. .16984
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L5767. .15789
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15184. .15321
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                                                                                                                                                                                                                                                                                                                                                                                            39.5%; Score 145.6; DB 9 96.7%; Pred. No. 1.6e-31;
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                                                                                                                                                                                                                                                                                        Worley, K.C
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                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished
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Rattus norvegicus
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Earnhart.C., Edgar,D., Edwards,C.C., Elhaj.C., Escotto,M., Falls,T., Ferraguto,D., Flagg,N., Ford,J., Foster,P., Erantz,P., Gabisi,A., Gao,J., Garcia,A., Garner,T., Garza,N., Gill,R., Garcial,J.H., Guevara,W., Gunaratne,P., Hale,S., Hamilton,K., Harris,C., Harris,K., Homes,M., Holloway,C., Hollins,B., Hernandez,O., Hodgson,A., Hogues,M., Holloway,C., Hollins,B., Homes,F., Howard,S., Huber,J., Hulyk,S., Hume,J., Jackson,L.E., Jackson,B., Jia,Y., Johnson,R., Jolivet,S., Joudah,S., Karlson,E., Kureshi,A., Landry,N., Leal,B., Lewis,L.C., Lewis,L., Li,J., Li,Z., Lichtarge,O., Lieu,C., Liu,J., Liu,W., Loulseged,H., Lozado,R.J., Lu,X., Lucier,R., Martindale,A., Martinez,E., Massey,E., Marhiney,E., McLeod,M.P., Meador,M., Mei,G., Metzker,M., Massey,E., Marhiney,E., McLeod,M.P., Meador,M., Mei,G., Metzker,M., Moser,M., Nufckerson,E., Newtson,N., Nguyen,A., Nguyen,N., Souton,S., Oguh,M., Okwuonu,G., Oragunye,N., Oviedo,R., Pace,A., Payton,B., Peery,J., Perez,L., Pitkens,R., Primus,B., Pu,L.L., Quiles,M., Ren,Y., Rives,M., Rojas,A., Rojubokan,I., Rolfe,M., Ruiz,S., Savery,G., Scherer,S., Scott,G., Shen,H., Shooshtari,N., Sisson,I., Sodergren,E., Sonaike,T., Sparks,A., Stanley,H., Stone,H., Suston,R., Wang,S., Ward-Moore,S., Warren,R., Washington,C., Watliamson,A., Wleczyk,R., Wooden,S., Watliamson,A., Wleczyk,R., Wooden,S., Watlings,G., Williamson,A., Wleczyk,R., Wooden,S., Watlings,G., Milliamson,A., Wleczyk,R., Wooden,S., Watlington,S., Williamson,A., Wleczyk,R., Wooden,S., Watlington,S., Williamson,A., Wleczyk,R., Wooden,S., Watlington,S., Williamson,A., Wleczyk,R., Wooden,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., Watlington,S., 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (13-JUL-2002) Human Genome Sequencing of Molecular and Human Genetics, Baylor College Baylor Plaza, Houston, TX 77030, USA On Jul 12, 2002 this sequence version replaced of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (11-FEB-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryan
Buhay, C., Burch, P., Burkett, C., Burrell, K. L., Byrd, N. C.,
Carron, T. F., Carter, M., Cavazos, S. R., Chacko, J., Chavez,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jul 12, 2002 this sequence version replaced g1:18847022
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Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
                                                                                                                                                                                          Center project name: GQLL Center clone name: CH230-140B23
                                                                                                                                                                                                                                                                                                    Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                    Web site: http://www.hgsc.bcm.tmc.edu/
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Center code: BCM
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NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 69 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
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REFERENCE AUTHORS

Rattus

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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

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Douthwaite, N. J., Draper, H., Dugan-Rocha, S., Durbin, N.J.,
Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M.,
Falls, T., Ferraguto, D., Flagg, N., Ford, J., Foster, P., Escotto, M.,
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Hernandez, D., Hodgson, A., Hogues, M., J.,
Jackson, E., Kally, S., Khan, U., King, L., Korvah, J., Kovar, C.,
Karlsson, E., Kally, S., Khan, U., King, L., Korvah, J., Lou, J., Liu, M., Liu, J., Liu, M., Lou, J., Liu, M., Lou, J., Liu, M., Lou, J., Liu, M., Lou, J., Liu, M., Lou, J., Liu, M., Lou, J., Liu, M., Lou, J., Liu, M., Lou, J., Liu, M., Martinez, E., Maseon, M., Mortis, S.,
Miller, G., Mitchell, T., Mohabbat, K., Morgan, M., Mortis, S.,
Noser, M., Nolkerson, J., Newtson, N., Nguyen, A., Nguyen, N.,
Nguyen, N., Nickerson, E., Newtson, N., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., Nguyen, A., N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (19-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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* NOTE: Estimated insert size may differ from sequence length

* (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft sequence. It currently

* consists of 69 contigs. The true order of the pieces

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as
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TITLE JOURNAL REFERENCE AUTHORS TITLE

Unpublished Direct Submission

(bases 1 to 190503)

JOURNAL

Direct Submission Worley, K.C.

Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap: version 0.990329
Consensus quality: 131119 bases at least 040
Consensus quality: 136808 bases at least 030
Consensus quality: 141839 bases at least 020

Web site: Center code: BCM

http://www.hgsc.bcm.tmc.edu/

Center: Baylor College of Medicine

runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.

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Isolated polypeptides associated with ovarian carcinomas, and the

WPI; 2001-211395/21.

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                                                                    14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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Best Local (
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                                                                                                                                                                                                                                                                       Ovarian
                                                                                                                                                                                                                                                                                                                     Ovarian carcinoma sequence
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10-SEP-1999;
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                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                  cancer; cancer therapy; vaccine; gene therapy; tumour; cancer;
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99US-0394374
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to ovarian tumour polynucleotides and polypeptides CC that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polypucleotides of the invention are CC useful for detecting a cancer in a pattent, for stimulating and/or CC expanding T-cells specific for a tumour protein, and for inhibiting the CC development of a cancer in a pattent. They are also useful for CC stimulating an immune response in a patient, and for treating a cancer in CC a patient and for determining the presence of a cancer in a patient. CC The isolated polynucleotides of the invention are useful for their CC ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and CC preparing ribozyme molecules for inhibiting expression of tumour CC polypeptides in tumour cells. Polypeptides and polynucleotides of the CC invention are also useful in recombinant DNA molecules to direct CC expression of a polypeptide in appropriate host cells. The sequences CC given in records ABL48760-ABL48956 represent polynucleotides encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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07-SEP-2000;
14-NOV-2000;
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(STOL/)
(ALGA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 369 BP; 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ovarian carcinoma proteins.
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                             CACCATCCT
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CACCATCCT
                                                                                         CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC
                                                                                                                          CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAA
                                                                                                                                                        CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
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) STOLK J A.
) ALGATE P A.
) FLING S P.
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2000US-0640173.
2000US-0656668.
2000US-0713550.
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01-DEC-2000;
20-DEC-2000;
20-DEC-2000;
20-DEC-2000;
28-FEB-2001;
10-MAY-2001;
25-MAY-2001;
                                                                   The invention relates to one hundred and twenty two nucleic acids encoding PRO polypeptides. The sequences of the 122 PRO polypnucleotides encode human secreted proteins. The PRO nucleic acids, polypeptides, agonists and antagonists are useful for treating a PRO related disorder. The PRO polypeptides are useful for diagnosing tumours, especially lung cancer, colon cancer, breast tumour, prostate tumour, rectal tumour or liver tumour. The PRO polypeptides are useful for stimulating the proliferation of, or gene expression, in pericyte cells, for stimulating the proliferation or differentiation of choolecyte cells, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; secreted protein; PRO; tumour; lung cancer; colon cancer; breast cancer; prostate tumour; rectal tumour; liver tumour; pericyte cell proliferation; chondrocyte cell proliferation;
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         stimulating the release of tumour necrosis factor alpha from human blood, for stimulating or inhibiting the proliferation of normal human dermal fibroblast cells. The PRO polypeptide may also be used as molecular weight markers and for tissue typing. The PRO nucleic acids have
                                                                                                                                                                                                                     Claim 2; Figure 15; 359pp; English.
                                                                                                                                                                                                                                                              One hundred and twenty two nucleic acids encoding PRO polypeptides, useful for treating a PRO related disorder and for diagnosing tumours such as lung cancer, colon cancer, breast tumour, prostate tumour, rectal
                                                                                                                                                                                                                                                                                                                                                                     Grimaldi JC,
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 applications in molecular biology,
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2001WO-US17092
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Smith V, Stephan JF,
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including use as hybridisation probes
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TF, Watanabe CK,
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Best Local :
     Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                    WPI;
                                                                         Хu J,
                                                                                                                   14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                   cytostatic; gene;
                                                                                                                                                                                                                                                             Human; ovarian cancer;
                                                                                                                                                                                                                                                                                Human ovarian carcinoma associated coding sequence SEQ ID NO: 205.
                                                                                                                                                                                                                                                                                                         05-SEP-2002
                                                                                                                                                                                                                                                                                                                                                   ABT03277 standard; cDNA; 1619
                                                                                              (CORI-) CORIXA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GGCAACTTTTTGCGGATTGTTCCTTGCTTNCAGGCTTTGCGCTGCAAATCCAGTGCTACCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in chromosome and gene mapping. {\tt ABK33536-ABK33657} represent human protein coding sequences of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGTGAACGTTCAAGACATGTGTCAGAAAGGAAGTGATGGAGCAAAGTGCCGGGATCATGTA
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                                                                        Stolk JA,
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56; Conservative
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                                                                         Algate
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99.2%;
                                                                                                                                                                                                                                                            ovarian carcinoma; gene therapy;
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Best Local Similarity
                                                                                                                                             14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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                                                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic; gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABT03281;
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                                                                                              (CORI-) CORIXA CORP
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                                                 Stolk JA,
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Best Local
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10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                         03-APR-2001; 2001US-0825294.
                                                                                                        US2002004491-A1
                                                                                                                                                                Ovarian cancer;
                                                                                                                                                                                     Ovarian carcinoma sequence isolate 57887
                                                                                                                                                                                                                                                           ABL40345 standard; cDNA; 1619 BP
                                                                                 10-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer
                                                                                                                                                                                                               28-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1619 BP; 399 A; 415 C; 432 G;
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                                                                                                                                                                                                                                                                                                                     CACCATCCT
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                                                                                                                                                                                                                                                                                                                                           CACCATCCT 369
                                                                                                                                                                                                                                                                                                                                                                              CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC
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99US-0394374.
2000US-0561778.
2000US-0640173.
2000US-0656668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                             (first entry)
                                                                                                                                                             cancer therapy; vaccine;
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Pred. No. 1.
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                                                                                                                                                            gene therapy; tumour;
                                                                                                                                                                                     extended cDNA.
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ABL40349 standard;

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CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT CCGCAAGTCCTGTGCATCAGCGGCCTGTCTCATCGCCCTCTGCCGGGTACCAGTCCTT

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GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGAAGCAAAGTGCCGGGATCATGTA 180

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GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGGAGCAAAGTGCCGGGATCATGTA

CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCAGGCCAGGGCTCCGCAC

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                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                          iseful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the extended cDNA sequence of ovarian carcinoma isolate 57887 given in record ABL48956.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (XUJJ/)
(STOL/)
(ALGA/)
                                                                                                                                                                                                                           Sequence 1619 BP; 399 A; 415 C; 432 G; 373 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (STOL/) STOLK J A. (ALGA/) ALGATE P A (FLIN/) FLING S P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
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GGCAACTTTTTGCGGATTGTTCTTGCTTNCAGGCTTTGCGCTGCAAATCCAGTGCTACCA 60
                                                                           GCAACTTTTTGCGGATTGTTCTTGCTTCCAGGCTTTGCGCTGCAAATCCAGTGCTACCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002-171027/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1a;
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                                                                                                                                                        Conservative
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                                                                                                                                                                     99.0%;
99.2%;
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                                                                                                                                                    Score 365.4; DB 24; Pred. No. 1.2e-109; 0; Mismatches 3;
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Best Local :
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                                                                                                                                                                                                                                                                                                               expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma 0591S nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2002004491-A1.
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                                                                                                                                                                                                                                                                                      Sequence 1619 BP; 399 A; 415 C; 432 G; 373 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim la; Page 125-126; 131pp; English.
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181 CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
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                                                                                                                                                                         1 GGCAACTTTTTGCGGATTGTTCTTGCTTNCAGGCTTTGCGCTGCAAATCCAGTGCTACCA 60
                               GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA
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                                                             GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 180
                                                                                                                              GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC 120
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STOLK J A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              detecting a cancer in a patient, for stimulating and/or
                                                                                                                                                                                                                          Conservative
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2000US-0656668.
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08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
protein/membrane protein expression. The nucleic acids and complementary sequences may also be used as DNA probes in diagnostic assays sequences may also be used as DNA probes in diagnostic assays can be called a complementary to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and activity. The antibodies may also be used as therapeutic agents to down regulate expression and activity. The antibodies may also
                                                                                                                                                                      This invention relates to nucleic acid sequences AAF93744 - AAF93916 which encode human secretory or membrane proteins represented by AAB88317 - AAB88419. Included in the invention are primers AAF93917 - AAF94295 and AAF62237 - AAF62235 which are used to isolate to CDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and diagnosis of diseases associated with inappropriate secretory.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-JUL-2000; 2000EP-0114090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; secretory protein; membrane protein; vaccine; gene therapy; rheumatoid arthritis; diabetes; ss.
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                                                                                                                                                                                                                                                                                                                                                                           Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy or as candidate target molecules
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                                                                                                                                                                                                                                                                                                                                                                           SEQ ID 203; 609pp + CD ROM; English
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2000JP-0183766
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                                                                                         14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
Novel ovarian cancer polypeptide and polynucleotide, useful detecting the presence of ovarian cancer in a patient, and i
                                 WPI; 2002-500186/53.
                                                      χu J,
                                                                                                                                                                                                                      Human; ovarian cancer; ovarian carcinoma; gene therapy; immunotherapy;
                                                                                                                                                                                                                                        Human ovarian carcinoma associated coding sequence SEQ ID NO:
                                                                        (CORI-) CORIXA CORP
                                                                                                                                                       23-MAY-2002.
                                                                                                                                                                           WO200239885-A2
                                                                                                                                                                                                                 cytostatic; gene;
                                                                                                                                   13-NOV-2001; 2001WO-US45395
                                                                                                                                                                                                                                                                 05-SEP-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             be used as diagnostic agents for detecting the presence of the polypeptides in samples (e.g. by enzyme linked immunosorbant assay (ELISA). Examples of diseases which may be treated include rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        arthritis and diabetes.
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D; Mismatches 3;
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                                                  Molesh DA;
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10-SEP-1999; 99US-0394374
01-MAY-2000; 2000US-0561778
                                                                             03-APR-2001; 2001US-0825294.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ovarian cancer; cancer therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ovarian carcinoma 01034C/0591S consensus nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-JUN-2002
                                                                                                                                          10-JAN-2002
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Pred. No. 1.2e-109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy; tumour; cancer;
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CC The invention relates to ovarian tumour polynucleotides and polypeptides concert therapy, be utilised in cancer therapy, for example in a vaccine or cc gene therapy. Polypeptides and polynucleotides of the invention are cuseful for detecting a cancer in a pattent, for stimulating and/or cepanding T-cells specific for a tumour protein, and for inhibiting the cepanding T-cells specific for a tumour protein, and for inhibiting the ct. a pattent of a cancer in a pattent. They are also useful for cc stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient cc ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and comparing ribozyme molecules for inhibiting expression of tumour cc polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current cc sequence represents the ovarian carrinoma offace comparing contents the current contents are completed in appropriate the current cc.
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07-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1897 BP; 435 A; 521 C; 532 G; 407 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence represents the ovarian carcinoma 01034C/0591S consensus
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                                                                                                                                                                                                                           CCGCAACTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
                                  CACCATCCT 369
                                                                                                    CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC
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CACCATCCT
                                                                  CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCAGGGCCAGGGCTCCGCAC
                                                                                                                                      CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 366;
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ALGATE P A.

FLING S P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stolk JA, Algate PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABB09417
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 127-128; 131pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; 2000US-0640173.
; 2000US-0656668.
; 2000US-0713550.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99.0%;
99.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 1.2e-109;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polynucleotide useful in diagnosis, cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1897;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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                                                                       639
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RESULT 12

Query Match Best Local Similarity

95.5**%**; 98.6**%**;

Score Pred.

352.4; DB 21; No. 2.4e-105;

Length 1953;

Sequence 1953 BP; 428 A; 540 C;

524 G;

458 T; 3 other;

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AAF22400/c
ID AAF22400 standard; cDNA; 1953 BP
                                                          CC ARMEGIBE. ARBGIBES to ARBGIZE represent more human secreted proteins are cally titles based on the tissues and cells the genes are expressed in. CE Examples of activities include: immunosuppressive; antiarthitic; CC examples of activities include: immunosuppressive; antiarthitic; CC examples of activities include: immunosuppressive; antiarthitic; CC exrebroprotective; nootropic; neuroprotective; antibacterial; virucide; CC exrebroprotective; nootropic; neuroprotective; antibacterial; virucide; CC exrebroprotective; nootropic; neuroprotective; antibacterial; virucide; CC exrebroprotective; nootropic; neuroprotective; antibacterial; virucide; CC exrebroprotective; nootropic; neuroprotective; antibacterial; virucide; CC exrebroprotective; nootropic; neuroprotective; antibacterial; virucide; CC exrebroprotective; nootropic; neuroprotective; antibacterial; virucide; CC exrebroprotection; norses, cats, dogs, chickens and proteins can be cC exist to prevent, treat or amedical condition in e.g. humans, cC expendights and pathological condition or susceptibility to a cC disorders e.g. confiction. Disorders which are diagnosed or treated include atomical anticity of the present of the breast or liver, cardiovascular CC disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral cC disorders e.g. corneal infection. The polypeptides can also be used to cald wound healing and epithelial cell proliferation, to prevent skin caping due to sunburn, to maintain organs before transplantation, for CC experience contaxis. The polypeptides can also be used as a food additive or cc preservative to increase or decrease storage capabilities. AAF22364 to AAF22372 and AAB63133 represent sequences used in the exemplification of the present interestion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fungicide; ophthalmological; gene therapy; autoimmune disease; neoplasm; rheumatoid arthritis; hyperproliferative disorder; cardiac arrest; rheumatoid arthritis; hyperproliferative disorder; cardiac arrest; cardiovascular disorder; cerebrovascular disorder; cerebral ischaemia; angiogenesis; nervous system disorder; Alzheimer's disease; infection; angiogenesis; nervous system disorder; Alzheimer's disease; infection; ocular disorder; corneal infection; wound healing; skin aging;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; secreted protein; diagnosis; immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF22400
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Isolated nucleic acid molecule encoding a human secreted protein used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-647420/62
P-PSDB; AAB63161.
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20-JAN-2000; 2000US-0176931.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              additive; preservative; ss.
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ROSEN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               preventing, treating or ameliorating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to AAF22421 encode the human secreted proteins given in AAB63134
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                                                  invention.
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       Claim 1; Page 443-444; 542pp; English
                                preventing, treating or ameliorating a disorder, Parkinson's diseases and cancers -
                                        Nucleic acid molecules encoding human secreted proteins, used in preventing, treating or ameliorating a disorder, e.g. Alzheimer's
                                                                                  P-PSDB;
                                                                                                                       Rosen CA,
                                                                                                                                                                                    11-JUN-1999;
                                                                                                                                                                                                            01-JUN-2000;
                                                                                                                                                                                                                                     21-DEC-2000
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                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted protein gene 13 SEQ ID NO:23
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DB; AAB75518.
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                                                                                                                                               ROSEN
                                                                                                                       Ruben SM,
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                                                                                                                                             GENOME SCI INC C A.
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                                                                                                                                                                                                                                                                                                                       disorder; skin ageing; chemotaxis;
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                             Human ovarian cancer associated coding sequence SEQ ID
                                                                                                                 AAF94818;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin ageing due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotraxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities. Included in the invention are polypucleotide sequences AAF64167 - AAF64175 and peptide AAB75505 which are used in the isolation, identification and characterisation of the proteins of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders
                                                                           23-MAY-2001
                                                                                                                                                      AAF94818 standard; cDNA; 396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1956 BP; 430 A; 543 C; 524 G; 456 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The proteins and polynucleotides, their agonists and antagonists have activities dependent on the tissues and cells in which they are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human secreted proteins AAB75506 - AAB75554 are encoded by polynucleotide sequences AAF64176 - AAF64224. The specification includes amino acid sequences AAB75555 - AAB75606 which represent fragments of the human secreted proteins, and protein sequences with which they share homology.
                                                                                                                                                                                                                                                                                                361
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                                                                                                                                                                                                                                                                                                CACCATCCT 369
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                                                                         (first entry)
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98.6%;
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                                                                                                                                                      ВP
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Pred. No. 2.4e-105;
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ovarian cancer; vaccine; gene therapy; carcinoma; ss

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated polypeptides associated with ovarian carcinomas, nucleic acids that encode them, useful for the prevention treatment of ovarian cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-SEP-1999; 99US-0394374
01-MAY-2000; 2000US-0561778
15-AUG-2000; 2000US-0640173
07-SEP-2000; 2000US-0656668
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 5; Page 119; 189pp; English.
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               ABT03085;
                                      ABT03085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAGCACCCCCTCTTTGTAA
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                                                                                                                                                                CGGGCCAAGGCCCAAGAAAAGGGGAA - - GTTCTGCCTCGGCCCTCANGCCATGGCTCCGC
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                                                                                                                                                  CGGGCCAAGGNCCAAAAAAAGGGGAAAGTTCTGNCCTCGGCCCTCAGGCCAGGGCTCCGC
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                                                                                                                         ACCACCATCCT 369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               396 BP; 90 A; 117 C; 95 G; 92 T; 2 other;
                                      standard;
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Pred. No. 1.1e-102;
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03-APR-2001;
02-OCT-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel ovarian cancer polypeptide and polynucleotide, useful detecting the presence of ovarian cancer in a patient, and i pharmaceutical compositions, e.g. vaccines, for treating over
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                                       CGGGCCAAGGNCCAAAAAAAGGGGAAAGTTCTGNCCTCGGCCCTCAGGCCAGGGCTCCGC
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359; Conserv
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                                                                                                                                                                                                                                                                                                                                         396
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; 2001US-0825294.
; 2001US-0970966.
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96.8%;
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completed:

November

7,

2002,

10:10:07

Job time : 78.6545 secs

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Maximum DB
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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4: /cgn2_6/ptodata/1,

5: /cgn2_6/ptodata/1,

6: /cgn2_6/ptodata/1,
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369
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1 US-08-035-392-31
1 US-08-592-126-97
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1 US-08-252-073A-5
2 US-08-9103-840A-1
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27.2	27.2	27.2	27.2	27.2	27.4	27.4	27.4	27.4	27.4	27.4	27.4	27.4	27.6	27.6	27.6	27.6	27.8
7.4	7.4	7.4	7.4	7.4	7.4	7.4	7.4	7.4	7.4	7.4	7.4	7.4	7.5	7.5	7.5	7.5	7.5
3596	3563	2348	1828	818	7803	7803	7705	7705	7705	7680	3487	656	3812	3355	344	343	8060
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US-08-779-801-5	US-09-041-886-20	US-09-393-554-10	US-08-858-207A-52	US-09-154-083-25	PCT-US93-12687-1	US-08-551-356-1	5455158-2	US-08-826-885-16	US-08-259-569-16	PCT-US95-09819-6	US-08-410-540-4	US-09-404-879A-313	US-09-784-316-1	US-08-991-944-3	US-09-423-233-26	US-09-423-233-27	US-08-766-528-1
	Sequence 20, Appl		Sequence 52, Appl	Sequence 25, Appl	Sequence 1, Appli	Sequence 1, Appli	Patent No. 5455158	Sequence 16, Appl	Sequence 16, Appl	Sequence 6, Appli	Sequence 4, Appli	Sequence 313, App	Sequence 1, Appli	Sequence 3, Appli	Sequence 26, Appl	Sequence 27, Appl	Sequence 1, Appli

## ALIGNMENTS

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; Sequence 6, Application ; Patent No. 5985655 ; GENERAL INFORMATION:
                                                                                              RESULT 2
US-08-484-126-6
                                                                                                                                                                                                                                                     Db 2688837 GCGGCTTGCGGGGCGCCAATGCGTTGGCCGTTGCGCCCGGGCGCCCCCTGGAACCCCGAAT 2688778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-103-840A-2
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                                                                                                                                                                         Db 2688717 ATCAGTGCAACCT 2688705
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Best Local S
Matches 71
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: FRASER, Claire M.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: DAN SEQUENCES FOR STRATITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: FLEISCHMAN, Robert APPLICANT: WHITE, Owen R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: CDC 1551
OTHER INFORMATION: "n" bases at
OTHER INFORMATION: represent a,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Mycobacterium tuberculosis
APPLICANT:
                                                                                                                                                                                                               323 GGAAGTTCTGCCT 335
                                                                                                                                                                                                                                                                                                                                                      203 GCGGCCTGTCTCATCGCCTCTGCCGGGTACCCAGTCCTTCTGCTCCCCAGGGAAACTGAAC 262
                                                                                                                                                                                                                                                                                        263 TCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAACGGGCCCAAGGCCCAAGAAAAGG 322
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Anderson, W. Baltrucki, L
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                                                                          US/08484126
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                                                                       RESULT 3
US-08-897-443-2/c
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       Sequence 2, Application US/08897443
Patent No. 5981263
GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 201-30-1
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
SEQUENCE TRACTERISTICS:
TENGTH: 759 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Lilite, Raymond J.78
REGISTRATION NUMBER: 31,778
REFERENCE/DOCKET NUMBER: 27:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,:
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Lillie, Raymond T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inc
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CORRECTORMORPHICES: 8
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                     123 CCTATCACCAATGCCACCCTGGACCAGC 150
                                                                                                                                                                                                                                                                                       220 CTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTG 279
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                                                                                                                                                                 340 CCTCANGCCATGGCTCCGCACCACCATC 367
                                                                                                                                                                                                                            280 CTGCAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAAGGGGAAGTTCTGCCTCGGC 339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/326,347
FILING DATE: 20-OCT-1994
APPLICATION NUMBER: 08/973,307
FILING DATE: 09-NOV-1992
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ADDRESSEE: Cecchi, Stewart & Olstein
STREET: 6 Becker Farm Road
                                                                                                                                                                                                                                                                                                                                      Local
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                                                                                                                                                                                              CTGAGCCTCCTCCTGCTGCATGCCCAGGACCCAGCGTGTGCCAACTTCTCGACCAGC
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507-513
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31,778
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RESULT 4
US-09-128-155-16/c
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GENERAL INFORMATION:
APPLICANT: Pan, Yang
APPLICANT: Pan, Yang
TITLE OF INVENTION: MOVEL MOLECULES OF TANGO-77 RELATED PROTEIN FAMILY
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: 09404/052001
CURRENT APPLICATION UNMBER: US/09/128,155
CURRENT FILING DATE: 1998-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 97; Conserv
                                                                                                                                                 Sequence 16, Application US/09128155 Patent No. 6117654
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLANCE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Billings Lucy J.
NAME: Billings Lucy J.
DE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Shah, Purvi
APPLICANT: Kaser, Mathew
TITLE OF INVENTION: HUMAN
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IMMEDIATE SOURCE:
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APPLICATION NUMBER:
FILING DATE:
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                                                                                                                                                                                                                                                                             458 ATGGTGCCCGTGGACAGATGCCGCATCCT 430
                                                                                                                                                                                                                                                                                                                 341 CTCANGCCATGGCTCCGCACCACCATCCT 369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        638 CCAATGGCAAAGATTAGGATGCCCGTGTCCCGTGCCTTAGCAGCCACCTCGGCCACGGAG 579
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                                                                                                                                                                                                                                                                                                                                                                                                281 TGCAACACCCCTCTTTGTAACGGGCCCAAGGCCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCC 340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    221 TCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGC 280
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CLONE: 681719
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/89 FILING DATE: Filed Herewith
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                                                                                                                                                                                                                                                                                                                                                            CGGGCCCCCTCTGCTTCTGAGAATGCGATGTTCAGGGCATACTGGATGGCCAGCCCGGTC
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3174 Porter Drive
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46.4%;
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                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17, Application US/09128155 Patent No. 6117654
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Best Local Similarity 49.4%;
Matches 79; Conservative
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 16
LENGTH: 152331
                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/128,155
CURRENT FILING DATE: 1998-08-03
EARLIER APPLICATION NUMBER: US 60/091,650
EARLIER FILING DATE: 1998-07-02
EARLIER APPLICATION NUMBER: US 60/054,646
EARLIER FILING DATE: 1997-08-04
NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Pan, Yang
TITLE OF INVENTION: NOVEL MOLECULES OF TANGO-77 RELATED PROTEIN FAMILY
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: 09404/052001
CURRENT APPLICATION NUMBER: US/09/128,155
                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(176373)
OTHER INFORMATION: n = A,T,C or
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EARLIER FILING DATE: 1998-07-02
EARLIER APPLICATION NUMBER: US 60/054,646
EARLIER FILING DATE: 1997-08-04
                                   77454 TTGGGGTCGGAGTACATTTGGGGATGAGGGGAGCTGTAAAAGCAGACTTGGACCTTTTCT 77513
                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (1)...(152331)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 176373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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156 TGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGC 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    156 TGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGC 195
                                                         36 TTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCT 95
                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         36 TTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCT 95
                                                                                                                                                                                                                                                                                                                                                                                                                                     FastSEQ for Windows Version 3.0
                                                                                                                                                                                           Conservative
                                                                                                                                                                                                          8.2%;
                                                                                                                                                                                       0; Mismatches 81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                          Score 30.4;
Pred. No. 23;
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Pred. No. 22;
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                                                                                                                                                                                                                          DB 3;
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                                                                                                                                                                                                                          Length 176373;
                                                                                                                                                                                       Indels
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                                                                                                                                                                                     0;
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                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                        RESULT 7
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                                                                                                                                                                                                                                                                               Patent No.
                                                                                                                                                                                                                                                                                                   Sequence 37,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/443,184A
CURRENT FILING DATE: 1999-11-19
NUMBER OF SEQ ID NOS: 138
SOFTWARE: PERL Program
SEQ ID NO 67
LENGTH: 327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                    APPLICANT: Brown, Michael S.
APPLICANT: Briggs, Michael R.
APPLICANT: Wang, Xiaodong
APPLICANT: Goldstein, Joseph I
TITLE OF INVENTION: METHODS AN
TITLE OF INVENTION: TO STEROL
TITLE OF INVENTION: TO STEROL
TITLE OF INVENTION: FROTEINS
NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Yue, Henry
APPLICANT: Baughn, Mariah
APPLICANT: Baimzai, Yalda
APPLICANT: Lal, Preeti
TITLE OF INVENTION: MAMMALIAN TOXICOLOGICAL RESPONSE MARKERS
FILE REFERENCE: PC-0007 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kaser, Matthew R. APPLICANT: Panzer, Scott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Cunningham,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Seilhammer, Jeffrey J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Zweiger, Gary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     77574 AGGAGCAGGAGGAGAAGAAGATGGATCTTAAGTGATTTGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: unsure LOCATION: 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: misc_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: a,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                         156 CCGTCCCTGGCTGTTGGACTCTTCTTTG 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               210 GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 269
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
                                                                                                                                                                                                                                                                                                                                                                                                                          270 GCATCAGCTGCTGCAACACCCCTCTTTG 297
                   STATE:
                                   CITY: Houston
                                                      STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            96 GGCTATGCGGCCCTGGTGGCTACTGGTGGGATTATTGNCTTTGCAAAAGCAGGTAGTGTG 155
COUNTRY: U.S.A.
                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    36 AAGTGACTGAACAATGCAGAAGGACAGTGGACCACTGGTTCCTTACATTATTATGGTTTC 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                 7, Application US/08131365B 5527690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                   Texas
                                                      P.O. Box 4433
                                                                                                                 GOLdstein, Joseph L.
VENTION: METHODS AND COMPOSITIONS RELATING
VENTION: TO STEROL REGULATORY ELEMENT BINDING
VENTION: PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                        Arnold,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Incyte ID No.
                                                                                                                                                                                                                   Michael R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        50.0%;
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                                                                        White & Durkee
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Pred. No. 1.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6372431 701649802H1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 3, Application US/08035392
Patent No. 5484732
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local
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TELEPHONE: (512) 418-3000
TELEFAX: (512) 474-7577
INFORMATION FOR SEQ ID NO: 37:
                                                                                                     CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: HOWells, Stacy L.
REGISTRATION NUMBER: 34,842
REFERENCE/DOCKET NUMBER: PD-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 455-5100
                                 TELEFAX: (619) 455-5110
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 2205 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/1:
FILING DATE: 01-OCT-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "DNA"
                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 19930322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Rosenfeld, M. G. APPLICANT: Andersen, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1771 AGATGGCCCTGGCTGGCCCAGTGGCTGCCCCC 1805
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                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    322 GGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCC 356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                262 CTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAACGGGCCAAGGCCCCAAGAAAAG 321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: CDS
LOCATION: 167..3607
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Parker, David L. REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 1880 Cent
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC COMPACTIVE OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                         90067
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               NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                         California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Spensley Horn Jubas & Lubitz
1880 Century Park East - Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                         USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TRANSCRIPTION FACTOR FOR THE REGULATION OF THE DEVELOPMENT OF SKIN AND HAIR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08/131,365
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Pred. No. 5.4;
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1771 AGATGGCCCTGGCTGGCCCAGTGGCTGCCTGCCCC 1805

322 GGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCC 356

US-08-668-123-37

Sequence 37, Application US/08668123 Patent No. 5891631

APPLICANT: Brown, Mich APPLICANT: Briggs, Mich APPLICANT: Wang, Xiaoo APPLICANT: Goldstein,

Wang, Xiaodong

INFORMATION: CANT: Brown, Michael S. CANT: Briggs, Michael R.

NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS: TITLE OF INVENTION:

PROTEINS

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

COUNTRY:

U.S.A.

STREET:

P.O. Box 4433

Arnold,

White & Durkee

Houston Texas

ADDRESSEE:

PRIOR APPLICATION DATA:

CLASSIFICATION:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/668,123 FILING DATE: 14-JUN-1996

US-08-131-365B-37

NAME/KEY: CDS LOCATION: 167..3607

FEATURE:

MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "DNA"

TOPOLOGY:

linear

STRANDEDNESS: single

nucleic acid

TELEFAX: (512) 474-7577
INFORMATION FOR SEQ ID NO: 37:
SEQUENCE CHARACTERISTICS:
LENGTH: 4154 base pairs

TELECOMMUNICATION INFORMATION: TELEPHONE: (512) 418-3000

REFERENCE/DOCKET NUMBER:

UTSD:372/PAR

ATTORNEY/AGENT INFORMATION:

TORNEY/AGENTAL DAVID L.
NAME: Parker, David L.
22,165

APPLICATION NUMBER: FILING DATE: 01-005 CLASSIFICATION: 43

01-OCT-1993 N: 435

Matches Query Match

Local Similarity

8.1%; 56.8%;

Score 30; DB Pred. No. 5.4; Mismatches

41;

Conservative

SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/131,365B

COMPUTER READABLE FORM:

MEDIUM TYPE:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

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                                                            US-08-504-511A-3
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             Query Match
Best Local Similarity
 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                   TELEPHONE: (619) 678-50
TELEFAX: (619) 678-5099
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
CITY: La Jolla
CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IMMEDIATE SOURCE:
CLONE: Skn-la
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: HOWells, Stacy L.
REGISTRATION UNMBER: 34,842
REFERENCE/DOCKET NUMBER: 07:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Rosenfeld, M. G. APPLICANT: Andersen, B. TITLE OF INVENTION: TRANSCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                     FEATURE:
                                                                                                                 CLONE: Skn-la
                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                             MOLECULE TYPE: CDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        245 TCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAACGGG 304
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                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: 20-JU CLASSIFICATION: 43
                                                                        NAME/KEY:
                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
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                                                                                                                                                                                                        LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      44;
 44;
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Conservative
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            8.0%;
64.7%;
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               Score 29.6; D
Pred. No. 5.4;
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Pred. No. 5.4;
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 Mismatches
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                             DB 1;
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 24;
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                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (415) 324-0880
TELEFAX: (415) 324-0960
INFORMATION FOR SEQ ID NO: 97:
                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
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TITLE OF INVENTION: Transcr:
TITLE OF INVENTION: Polypepi
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 46
TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: US
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                                                                                                                                                                                                                                                                                                                                                            ORIGINAL SOURCE:
INDIVIDUAL ISOLATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Sholtz, Charles K.
REGISTRATION NUMBER: 38,
                                                                                                                                                                                                          1392 GCGTTGGTCTCCTCCTCCAGTTCCCGGCGCTTTTCCTCCACCTTGCGCTTCTCCTCCTGG 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
1212 AGCTCACTTAGGAACTCCTTCCT 1190
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CTTY: Palo Alto
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                                                                                                                                                                                                                                        167 GCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCC 226
                                  347 CCATGGCTCCGCACCACCATCCT 369
                                                                                                   287 ACCCCTCTTTGTAACGGGCCCAAGGCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCCCTCANG
                                                                                                                                                           227 GGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTTCAGTTTGCATCAGCTGCTGCAAC 286
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  32 CCAAAGCC 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
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                                                                                                                                     TGGACCCGCTTCAGGTGCTCAAACTTCTCATGGAGCTCCCTTTCCTTCTCCTTCAGCTCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1534 base pairs
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Pred. No. 6.1;
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US-08-292-945-5
Sequence 5, Application US/08252073A Patent No. 5767073
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Patent No. 558547
GENERAL INFORMAT
                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                          Matches
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INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
FILING DATE: December 10,
ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,28
                                                                                                                                                                                                                                                                                                                                                                                                         ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                           HYPOTHETICAL: I
                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: ME
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404) 815-6508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                     405
                                                                                                                                                                                            128 GTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAG 187
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                                                                                              465 TATGGGCCCCGACCAGAGGAGTACGAATTCCTCACTCCAG 504
                                                                                                                              188 TCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCG 227
                                                                                                                                                                                                                                345 GAATACAGGGTGAAAATTAACTTCAAAGTGAATAAGGATATTGTGTCTGGCCTGAAGTAT 404
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                                                                                                                                                                                                                                                                                                                         Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/292,945
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 1100
CITY: Atlanta
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                                                                                                                                                              GTTCAACACACATACCGGACTGGCATGAGAGTGGATAAAGCCACATTCATGGTTGGCAGC 464
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78; Conservative
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1100 Peachtree Street, Suite 2800
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Adra, Chaker
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SYSTEM: PC-DOS/MS-DOS
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48.8%;
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                                                                                                                                                                 Sequence 5, Application PC/TUS9312074 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
CURRENT APPLICATION DATA:
                                                                                     TITLE OF INVENTION: D4 Gene and Methods of Use Thereof NUMBER OF SEQUENCES: 8
COMPUTER READABLE FORM:
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LENGTH: 1107 base pairs
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APPLICATION NUMBER: US 07/990,337
FILING DATE: 10-DEC-1992
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                              465 TATGGGCCCCGACCAGAGGAGTACGAATTCCTCACTCCAG
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                                                                                                                                                                                                                                                                                                               188 TCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCG 227
                                                                                                                                                                                                                                                                                                                                                  405 GTTCAACACACATACCGGACTGGCATGAGAGTGGATAAAGCCCACATTCATGGTTGGCAGC 464
                                                                                                                                                                                                                                                                                                                                                                                       128 GTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAG 187
                                                                                                                                                                                                                                                                                                                                                                                                                        345 GAATACAGGGTGAAAATTAACTTCAAAGTGAATAAGGATATTGTGTCTGGCCTGAAGTAT 404
             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0
                                                                       MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              68 GAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAAC 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: ME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center,
STREET: 1201 West Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM:
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SYSTEM: PC-DOS/MS-DOS
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48.8%;
             Release #1.0, Version #1.25 (EPO)
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D4 Gene and Methods of Use Thereof
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 Mismatches

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APPLICATION NUMBER: US 07/9
FILING DATE: 10-DEC-1992
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1107 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear

MOLECULE TYPE: CI

CDNA

APPLICATION NUMBER: PCT/US93/12V...
FILING DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/990,337

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Sequence 2. Application US/09103840A
Patent No. 6294328
GENERAL INFORMATION:
APPLICANT: FLEISCHMAN, Robert D.
APPLICANT: WHITE, Owen R.
APPLICANT: VENTER, John C.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: TUBERCULOSIS
FILE OF INVENTION: TUBERCULOSIS
FILE OF INVENTION UMBER: US/09/103,840A
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
TYPE: DNA
CONTRACT AND CONTRACTOR OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
TYPE: DNA
CONTRACTOR OF SEQ ID NOS: 2
TYPE: DNA
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Matches 42; Conserv
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Best Local Similarity 48.8%;
Matches 78; Conservative
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FEARURE:
OTHER INFORMATION: CDC 1551
OTHER INFORMATION: "n" bases at various positions throughout the sequence
OTHER INFORMATION: represent a, t, c or g
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ORIGINAL SOURCE:
ORGANISM: Muri
                                                                              359 ACCAC 363
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Search completed: November 7, 2002, 15:58:03 Job time : 3169.83 secs

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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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      Score
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/cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
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ch 99.5%; Score 367; DB 10; Length 369; L Similarity 100.0%; Pred. No. 1.6e-111; 369; Conservative 0; Mismatches 0; Indels 0; GGGAGTTTTTTGCGGGATTGTTGTTGCTTGCTTCTTGCTTTGCGCTGCAAATCCAATCCAGTGCTACCA	/09825294 TIONS AND METHODS FOR THE Y AND DIAGNOSIS OF OVARIAN CANCE US/09/825,294 04-03 WS Version 3.0	US-09-864-761-4612 US-09-780-525-1 US-09-780-525-1 US-09-923-876-5109 US-09-960-352-14418 US-09-764-853-72 US-09-867-701-5083 US-09-867-701-5284 ALIGNMENTS	US-09-764-846-136 US-09-764-846-136 US-09-822-830A-187 US-09-934-060A-23 US-09-934-060A-29 US-09-934-060A-3 US-09-934-060A-1 US-09-934-060A-3 US-09-934-060A-3 US-09-934-060A-3 US-09-934-060A-3 US-09-934-060A-3 US-09-940-060A-3 US-09-940-060A-3 US-09-956-74-30-6410 US-09-216-393-97 US-09-974-300-6410 US-09-905-291A-33 US-09-905-291A-33 US-09-909-320-33 US-09-909-320-33 US-09-909-320-33 US-09-909-320-33 US-09-909-320-33 US-09-909-320-33 US-09-909-320-33 US-09-909-320-33
Length 369; ; Indels 0; Gaps 0; CAAATCCAGTGCTACCA 60 [	ANCER .	Sequence 4612, Ap Sequence 1, Appli Sequence 5109, Ap Sequence 14418, A Sequence 72, Appl Sequence 5083, Ap Sequence 5083, Ap Sequence 5284, Ap	Sequence 66, Appl Sequence 136, App Sequence 23, Appl Sequence 29, Appl Sequence 5, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1754, Ap Sequence 1754, Ap Sequence 964, Appl Sequence 97, Appl Sequence 97, Appl Sequence 33, Appl Sequence 33, Appl Sequence 33, Appl Sequence 33, Appl Sequence 21354, Ap Sequence 21354, Appl Sequence 21354, Appl

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APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Filng, Steven P.
APPLICANT: Filng, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 21011.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 205
LENGTH: 1619
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RESULT 3
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Matches 366;
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Local Similarity 99.2%;
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Pred. No. 1.1e-110;
0; Mismatches 3;
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GENERAL INCOMPATION

GENERAL INCOMPATION

APPLICANT: Xu, Jiangchun

APPLICANT: Stolk, John A.

APPLICANN: Algate, Paul A.

APPLICANN: Algate, Paul A.

APPLICANN: Algate, Paul A.

APPLICANN: Fing, Steven P.

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA

FILE REFERENCE: 20111.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTMARE: FrastSEQ for Windows Version 3.0

SEQ ID NO 211

LENGTH: 1619

TYPE: DNA

ORGANISM: Homo sapiens
Sequence 214, Application US/09825294
Patent NO. US20020004491A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Hing, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT APPLICATION AND FOR THE COURTENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 214
LENGTH: 1897
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99.2%;
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                                                                                                                                                   OF OVARIAN CANCER
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GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Algate, Paul A.

APPLICANT: Filng, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 9

LENGTH: 396
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; OTHER INFORMA
US-09-825-294-214
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                                                                                                               ; NAME/KEY: misc_feature; LCCATION: (1)...(396)
; OTHER INFORMATION: n = A,T,C
US-09-825-294-9
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US-09-825-294-9
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                                                       Matches 359;
                                                                                   Query Match
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FEATURE:
NAME/KEY: misc_feature
LCCATION: (1)...(1897)
OTHER INFORMATION: n = A,T,C
                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
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Pred. No. 9.7e-104;
0; Mismatches 10;
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Pred. No. 1.2e-110;
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APPLICANT: XLO, Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Algate, Paul A.

APPLICANT: Algate, Paul A.

APPLICANT: Fling, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN

FILE REFERENCE: 210121, 484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03
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; Patent No. US20020004491A1
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NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 212

LEMONT: 222
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                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 1010
TYPE: DNA
 300
                              467
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A-CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGC
                                            TCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTA 299
                                                                                                      ACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTTGCCGGGTACCAGTCCT
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                                                                                                                                                                                                                                                                                                                                   Score 341.8; DB 10;
Pred. No. 5.2e-103;
0; Mismatches 4;
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                                   ; ORGANISM: Homo sapien 
US-09-867-701-4251
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US-09-867-701-4251
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US-09-825-294-213
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LENGTH: 430
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SOFTWARE: FRATSEQ for Windows Version 3.0
SEQ ID NO 213
LENGTH: 480
    Query Match
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                                                                                                                                       APPLICANT: AGLATE, Paul A.
APPLICANT: AGLATE, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. US20020004491A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN
FILE REFERENCE: 210121.484C5
                                                                                                               NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
                                                               TYPE: DNA
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Pred. No. 2.3e-55;
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Length 430;
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AND

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APPLICANT: Wasten, Wesley C.
APPLICANT: Tao, Nenghing
APPLICANT: Byatt, John C.
APPLICANT: Byatt, John C.
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
FILE REFERENCE: 16511.006/37-21(10298)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
SEQ ID NO 3015
SEQ ID NO 3015
LENGTH: 407
TYPE: DNA
ORGANISM: Bos taurus
OTHER INFORMATION: Clone ID: 13-LIB34-071-Q1-E1-D1
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APPLICANT: Warren, Wesley C.
APPLICANT: Tao, Nengbing
APPLICANT: Tao, Nengbing
APPLICANT: Byatt, John C.
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REPERENCE: 16511.006/37-21(10298)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
SEQ ID NO 13056
LENGTH: 412
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Patent No. US20020137139A1
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74; Conservative
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; OTHER INFORMATION: unsure at all n locations
; OTHER INFORMATION: Clone ID: 56-LIB188-023-Q1-E1-F8
US-09-960-352-13056
                                                                                                                                                              RESULT 12
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                                                              APPLICANT: Warren, Wesley C.
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APPLICANT: Mathialagan, Nagappan
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION.
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REFERENCE: 16511.006/37-21(10298)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
SEQ ID NO 9456
LENGTH: 423
                                                                                                  Sequence 1886, Application US/09960352 Patent No. US20020137139A1
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Best Local Similarity
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           APPLICANT:
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ORGANISM: BOS taurus
OTHER INFORMATION: Clone ID: 41-LIB188-009-Q1-E1-C10
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Tao, Nengbing
Byatt, John C.
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Pred. No. 0.3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Bolt, Andrew
APPLICANT: Van Huffel, Christophe
APPLICANT: Van Huffel, Christophe
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE IDENTIFICATION, ASSESSMENT, F
TITLE OF INVENTION: OF HUMAN CANCERS
FILE REFERENCE: MRI-016B
CURRENT APPLICATION NUMBER: US/09/834,975
CURRENT FILING DATE: 2001-04-13
PRIOR FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 60/197,538
PRIOR FILING DATE: 2000-04-14
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Best Local Similarity 51.4
74; Conservative
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SEQ ID NO 1886
LENGTH: 458
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ORGANISM: Homo sapiens
FEATURE:
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235 GTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCT
                                             144 TGCATACAATTGGTTTTNNGAATNANAACTGGTCAATTTTTCTGCCTATGNGGGCTACTT
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                                                                                                                                           84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      84
                                                                                                                                                                                                                                                             Local Similarity
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                                                                                         CATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCA 234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACGACTGCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTC 143
                                                                                                                                       TTGCATGGTNCCTTTTCAACTCATGGGCCACAGGNGACTCTTANAGATTTATGCCAAAAT 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGGCCAACGACCTGTTTGAAAGTG 325
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                                                                                                                                                                                                                                                      Score 31.6; DB Pred. No. 0.73;
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204 TCAGTTTNGTTCTNAATNAACATTTTNGACTTTAANAAGAGCCTNCATTTGCCCCCTTTNT 263

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RESULT 15
US-10-03-109-13
Sequence 13, Application US/10033109
Patent No. US20020142390A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Allen, Stephen M.
APPLICANT: Rafalski, J. Antoni
APPLICANT: Sakal, Hajime
TITLE OF INVENTION: Nitrogen Transport Metabolism
FILE REFERENCE: BB-1210
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; ORGANISM: Arabidopsis thaliana
US-09-770-445-893
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US-09-770-445-893
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SEQ ID NO 893
LENGTH: 775
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Expressed Sequences of Arabidopsis TITLE OF INVENTION: thaliana FILE REFERENCE: 2023US (PARA-012PRV) CURRENT APPLICATION NUMBER: US/09/770,445 CURRENT FILING DATE: 2001-01-26 PRIOR APPLICATION NUMBER: US 60/178,472 PRIOR FILING DATE: 2000-01-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
CURRENT APPLICATION NUMBER: US/10/033,109
CURRENT FILING DATE: 2001-12-28
PRIOR APPLICATION NUMBER: BARLIER APPLICATION NUMBER: 09/384,625
PRIOR FILING DATE: EARLIER FILING DATE: 1999-08-27
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                                                                                                                                                                                                                                                                                                                     149 GAAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCC 190
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                                                                                                                                                                                                                                                                                              409 GGAGTGGAGAAGGGAACGGCGGAGCTAAGGCCCAACCAAGTAC 450
                                                                                                                                                                                                                                                                                                                                                                       349 TTCTCCTCTGACGATTTCATTGGCGAAGCCACGGTACATGTGAAAGAGCTATTGGAAATG 408
                                                                                                                                                                                                                                                                                                                                                                                          89 TGCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAA 148
                                                                                                                                                                                                                                                                                                                                                                                                                                    8.6%;
Local Similarity 56.9%;
es 58; Conservation
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Hamilton, Carol M.
Price, Jennifer L.
Raines, Tracy M.
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Rameaka, Joshua G.
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Hurban, Patrick
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; TYPE: DNA
; ORGANISM: Triticum aestivum
US-10-033-109-13
Search completed: November Job time: 19.0364 secs
                                                                                      밁
                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/098,248
PRIOR FILING DATE: EARLIER FILING DATE: 28 August 1998
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Microsoft Office 97
SEQ ID. 0 13
                                                                                                                                                                                                                                                           Best Local Similarity
Matches 62; Conserv
                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                            266 GTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAACGGGCCCAAGGCCCCAAGAA 318
                                                                                                                                                                       645 GGCTTCCTCCTCCACTGGGGCGTCATCGACTACTGCGGCGGCTACGTCATCCACATCCCC 704
                                                                                   705 GCCGGCGTCGCCGGCTTCACCGCCGCGTACTGGGTCGGGCCAAGGACCAAGAA 757
                                                                                                                                                                                                                 206 GCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCA 265
                                                                                                                                                                                                                                                           Conservative
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                          2002, 17:59:58
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pred. No. 1.5;
0; Mismatches 51;
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Database
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Maximum Match 100%
Listing first 45 s
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Maximum DB seq length: 200000000
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Perfect score:
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369
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Copyright (c) 1993 - 2002 Compugen Ltd.
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11220.326 Million cell updates/sec
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76: /cgn2_6/ptodata/1/pna/US6012_COMB.seq:
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78: /cgn2_6/ptodata/1/pna/US6013_COMB.seq:
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79: /cgn2_6/ptodata/1/pna/US6013_COMB.seq:
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79: /cgn2_6/ptodata/1/pna/US6013_COMB.seq:
79: /cgn2_6/ptodata/1/pna/US6013_COMB.seq:
79: /cgn2_6/ptodata/1/p
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.

Score

Match Length DB

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1524	1524	1524	1524	1524	1524	1524	1524	1524	1524	1524	1524	1524	1524	1524	369	369	369	369	369	369
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US-10-218-849-15	US-10-218-784-15	US-10-218-765-15	US-10-218-631-15	US-10-218-612-15	US-10-216-168-15	US-10-216-167-15	US-10-216-166-15	US-10-216-165-15	US-10-216-164-15	US-10-216-163-15	US-10-216-162-15	US-10-216-160-15	US-10-216-159A-15	US-10-119-480-15	US-10-212-677-199	US-09-970-966-199	US-09-825-294-199	US-09-713-550-199	US-09-656-668-199	PCT-US01-45395-199
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APPLICANT: Stolk, John A.

APPLICANT: Algate, Paul A.

APPLICANT: Filing, Steven P.

APPLICANT: Molesh, David Alan

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.48401PC

CURRENT APPLICATION NUMBER: PCT/US01/45395

CURRENT FILING DATE: 2001-11-13

NUMBER OF SEQ ID NOS: 215

SOFTMARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 199

LENGTH: 369
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PCT-US01-45395-199
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GENERAL INFORMATION:
APPLICANT: Corixa Corporation
APPLICANT: Xu, Jiangchun
                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION: 29, 345
OTHER INFORMATION: n =
                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens FEATURE:
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                       GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
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US-10-219-010-15
US-10-219-061-15
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US-10-219-064-15
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SEQ ID NO 199
LENGTH: 369
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(369)
OTHER INFORMATION: n = A,T,C or
US-09-656-668-199
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APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
TITLE OF INVENTION: OVARIAN TUMOR SEQUENCES
TITLE OF INVENTION: METHODS OF USE THEREFOR
FILE REFERENCE: 210121.484C3
CURRENT APPLICATION NUMBER: US/09/656,668
CURRENT FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 199
SOFTWARE: FastSEQ for Windows Version 3.0
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369; Conserv
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                                                                       369
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OF USE THEREFOR
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Sequence 199, Application US/09825294

GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Xtolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Filing, Steven P.
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION UNUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
INUMBER OF SEQ ID NOS: 215
SOFTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 199
LENGTH: 369
TYPE: DNA
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Sequence 199, Application US/09713550
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Xtolk, John A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C4
CURRENT APPLICATION UNMBER: US/09/713,550
CURRENT FILING DATE: 2000-11-14
NUMBER OF SEQ ID NOS: 205
COPTMANDE: PARTSFOR FOR WINDOWS VERSION 3.0
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LOCATION: (1)...(369)
OTHER INFORMATION: n = A,T,C or
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ORGANISM: Homo sapiens
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APPLICANT: Fling, Steven F.
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
TITLE REFERENCE: 210.121.484C6
CURRENT APPLICATION NUMBER: US/09/970,966
CURRENT ETLING DATE: 2001-10-02
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
LENGTH: 369
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                                                                                                      NAME/KEY: misc_feature; LOCATION: 29, 345; OTHER INFORMATION: n = A,T,C or GUS-09-970-966-199
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Best Local Similarity
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                                                 Matches 369;
                                                                             Query Match
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Molesh, David Al.
Fling, Steven P.
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                                                   Conservative
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                                                                99.5%; Score 367; 100.0%; Pred. No.
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                                                                                 DB 36;
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FEATURE:
NAME/KEY: misc_feature
LOCATION: 29, 345
OTHER INFORMATION: n = A,T,C or G
US-10-212-677-199
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CURRENT FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 288
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
LENGTH: 369
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GENERAL INFORMATION
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APPLICANT: Xu, Jiangchun
APPLICANT: Fanger, Gary R.
APPLICANT: Harlocker, Susan L.
APPLICANT: Herlocker, Susan L.
APPLICANT: McNeill, Patricia D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIACROSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C7
                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
241 CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAA 300
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                                                                       CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT 240
                                                                                                GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
                                                                                                                                                                       GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC 120
                                           GGCAACTTTTTGCGGATTGTTCTTGCTTNCAGGCTTTGCGCTGCAAATCCAGTGCTACCA 60
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 Mismatches

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US-10-119-480-15
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LENGTH: 1524
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/119,480 CURRENT FILING DATE: 2002-04-09 NUMBER OF SEQ ID NOS: 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Baker, Kevin P. APPLICANT: Desnoyers, Luc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo Sapien
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501 CACCATCCT
                                              361 CACCATCCT 369
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Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wood, William I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Stephan, Jean-Philippe F. Watanabe, Colin L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Desnoyers, Luc Gerritsen, Mary
                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                              99.0%; Score 365.4; DB 4 99.2%; Pred. No. 3.3e-98;
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RESULT 8 US-10-216-159A-15

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FILLE OF INVESTION. OCCUPANTS.
FILL REFERENCE: P3550PIC6

CURRENT APPLICATION NUMBER: US/10/216,159A

CURRENT FILING DATE: 2002-08-09

PRIOR APPLICATION NUMBER: 10/119,480

PRIOR FILING DATE: 2002-04-09

PRIOR APPLICATION NUMBER: 60/059113

PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/062287

PRIOR APPLICATION NUMBER: 60/06287

PRIOR FILING DATE: 1997-10-28

PRIOR FILING DATE: 1997-10-31

PRIOR APPLICATION NUMBER: 60/064103

PRIOR FILING DATE: 1997-10-31

PRIOR APPLICATION NUMBER: 60/064103

PRIOR FILING DATE: 1997-10-31

PRIOR APPLICATION NUMBER: 60/064973

PRIOR APPLICATION NUMBER: 60/069873
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PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR ETILING DATE: 1998-03-25
PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079656
PRIOR FILING DATE: 1998-03-26
PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079728
PRIOR APPLICATION NUMBER: 60/079728
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Baker, Kevin P.
APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo Sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 1524
                                                                                                                                                                                                                                                                                                                                                                                121
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                                                                         381 CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                         GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCCCCCGAGTTCATTGTGAATTGCAC 120
                                                                                                                                                                                                                               GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 180
                                    CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC 360
CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA 300
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Gerritsen, Mary
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Smith, Victoria
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Grimaldi, J. Christopher
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99.0%; Score 365.4; DB 42; 99.2%; Pred. No. 3.3e-98;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3; Indels
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US-10-216-160-15
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NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
US-10-216-162-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: P3530P1C8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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                                                                                                                                             301 CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC 360
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                                                           501 CACCATCCT 509
                                                                                      361 CACCATCCT 369
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                                                                                                                                                                         CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAA 300
                                                                                                                                                                                                                                                                                              CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
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Gerritsen, Mary
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Grimaldi, J. Christopher
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99.0%; Score 365.4; DB 42; Length 1524; 99.2%; Pred. No. 3.3e-98; tive 0; Mismatches 3; Indels 0;
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Sequence 15, Application US/10216162 GENERAL INFORMATION: APPLICANT: Baker, Kevin P. APPLICANT: Desnoyers, Luc

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CURRENT ETLING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/19,480
PRIOR FILLING DATE: 2002-04-09
PRIOR FILLING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILLING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/0709294
PRIOR APPLICATION NUMBER: 60/0709294
PRIOR APPLICATION NUMBER: 60/0709294
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PRIOR APPLICATION NUMBER: 60/0709294
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SEQ ID NO 15
LENGTH: 1524
TYPE: DNA
ORGANISM: Homo Sapien
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APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC FILE REFERENCE: P3530P1C2
501 CACCATCCT 509
                                        361 CACCATCCT 369
                                                                                                                              301
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                                                                                                                                                                                                                                                                               181 CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT 240
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REPLICATION NUMBER: 60/079728

REPLICATION NUMBER: 60/079728

REPLICATION NUMBER: 60/079728
                                                                 CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCAGGCCAGGGCTCCCGCAC
                                                                                        CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC 360
                                                                                                                                                                                                                                       GETGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGGATCATGTA
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Grimaldi, J. Christopher
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99.2%;
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US-10-216-163-15
; Sequence 15, Applica
; GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: 10/119/480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/05287
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-17
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
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PRIOR APPLICATION NUMBER: 60/079656
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APPLICANT: Desnoyers, Luc
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PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC FILE REFERENCE: P3530P1C3
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381 CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA 440
                   241 CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCTGCAACACCCCTCTTTGTAA 300
                                                                                                                                     181
                                                                                                                                                                               121 GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 180
                                                                                                                                                            261 GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 320
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                                                                            CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT 380
                                                                                                   CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT 240
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Grimaldi, J. Christopher
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Gerritsen, Mary
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 Mismatches

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PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079656
PRIOR APPLICATION NUMBER: 60/079656
                                                                                                                                                                                               Query Match
Best Local Similarity 99.2
                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 15
LENGTH: 1524
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PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/216,164
CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                       Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 246
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                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1998-03-27
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APPLICANT: Desnoyers, I
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PRIOR FILING DATE: 1997-10-31
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                               121 GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 180
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                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/079728
                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 1998-03-26
GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA
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Grimaldi, J. Christopher
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99.2%;
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                                                                                                                                                                                                                               Score 365.4; DB 42; Length 1524; Pred. No. 3.3e-98;
                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                              Indels
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PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
                                                                                                                                                                                         Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 246 SEQ ID NO 15 LENGTH: 1524
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                                                                      Matches 366;
                                                                                                      Query Match
                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1998-03-27
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PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
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APPLICANT: Desnoyers, Luc
                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/079656
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PRIOR FILING DATE: 1998-03-25
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                                                                                                                                                     TYPE: DNA
ORGANISM: Homo Sapien
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                1 GGCAACTTTTTGCGGATTGTTCTTGCTTNCAGGCTTTGCGCTGCAAATCCAGTGCTACCA 60
                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/078910
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GGCAACTTTTTGCGGATTGTTCTTGCTTCCAGGCTTTGCGCCTGCAAATCCAGTGCTACCA 200
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FILING DATE: 2002-08-09
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Gerritsen, Mary
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Grimaldi, J. Christopher
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                                                                      Conservative
                                                                                  99.0%;
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                                                                    0,
                                                                                  Score 365.4; DB Pred. No. 3.3e-98
                                                                  Mismatches
                                                                                                                                                                                                                                        See File Wrapper or PALM.
                                                                                                  DB 42;
                                                                    Indels
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US-10-216-166-15
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                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/069873
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
PRIOR FILING DATE: 1998-03-25
                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/216,166
CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 00/062287
PRIOR APPLICATION NUMBER: 00/062287
PRIOR APPLICATION NUMBER: 00/062287
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APPLICANT: Desnoyy
APPLICANT: Gerrit:
APPLICANT: Goddar
APPLICANT: Goddws
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APPLICANT: Grimal
APPLICANT: Smith,
APPLICANT: Stephal
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GENERAL INFORMATION:
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APPLICANT: Wood, William I.

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: 93530PIC9
                                                                                                Remaining Prior P
                                                                                                                                        PRIOR FILING DATE: 1998-03-27
                TYPE: DNA
ORGANISM: Homo
                                                          LENGTH:
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                                                                         NO 15
                                                                                                                                                                             APPLICATION NUMBER: 60/079656 FILING DATE: 1998-03-26
                                                                                                                                                           APPLICATION NUMBER: 60/079728
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                                                          1524
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Gerritsen, Mary
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                                                                                                Application data removed - D NOS: 246
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                                                                                                                   See File Wrapper or PALM
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APPLICANT: SCEPHEN, SCENT FILIPPE F.
APPLICANT: WACANABA, COLIN L.
APPLICANT: WOOD, WILLIAM I.
FITTLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FITTLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C4
CURRENT APPLICATION NUMBER: US/10/216,167
CURRENT APPLICATION NUMBER: US/10/216,167
CURRENT FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-10-18
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/069873
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
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Smith, Victoria
Stephan, Jean-Philippe F.
Watanabe, Colin L.
Wood, William I.
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Gerritsen, Mary
Goddard, Audrey
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NUMBER OF SEQ ID NOS: 246
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LENGTH: 1524
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Application US/10230437 WATION: Baker, Kevin P. Besnoyers, Luc Gerritsen, Mary Goddard, Audrey Godowski, Paul J.	ALIGNMENTS	6 US-10-240-425-364	5 US-09-620-607B-20	US-09		-00-SD	US-10-240	-01-Sn	US-10-	US-10		US-10	US-10	us-10-128-685 <i>t</i>	US-10-1	US-10-127-849A-	6 US-10-127-843A-293	
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; ORGANISM: HOMO US-10-230-437-15
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PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/079910
PRIOR FILING DATE: 1998-03-20
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079294
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PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-26
PRIOR FILING DATE: 1998-03-27
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Query Match
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Gurney, Austin L.
Smith, Victoria
Stephan, Jean-Philippe F
Watanabe, Colin L.
Wood, William I.
                                                                                                                                                                                                                            Application
                                                                                  Sapien
99.0%;
                                                                                                                                                                                                                          data removed -
Score 365.4;
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DB
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Length
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Best Local Similarity

99.2%;

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; ORGANISM: Homo sapiens US-10-085-198-105
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                                                                                                                           SEQ ID NO 105
LENGTH: 4321
                                     Query Match
       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/085,198
CURRENT FILING DATE: 2002-02-25
                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: 60/277,324
                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/312,858
PRIOR FILING DATE: 2001-08-16
PRIOR APPLICATION NUMBER: 60/271,840
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/311,981 PRIOR FILING DATE: 2001-08-13
                                                                                                                                                                           Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 653
                                                                                                                                                                                                                PRIOR FILING DATE: 2001-02-28
                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/272,405
                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/315,614 PRIOR FILING DATE: 2001-08-29
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PRIOR APPLICATION NUMBER: 60/299,695
PRIOR FILING DATE: 2001-06-20
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                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2001-03-20
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PRIOR FILING DATE: 2001-02-26
Local 5.71;
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                     Similarity
                                                                                                                                                             PatentIn Ver.
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   8.5%;
ilarity 51.4%;
Conservative
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                  Score 31.4;
Pred. No. 1
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PRIOR APPLICATION NUMBER: 60/271,646
PRIOR APPLICATION NUMBER: 60/276,401
PRIOR APPLICATION NUMBER: 60/276,401
PRIOR APPLICATION NUMBER: 60/276,401
PRIOR APPLICATION NUMBER: 60/311,981
PRIOR FILING DATE: 2001-08-13
PRIOR APPLICATION NUMBER: 60/312,858
PRIOR APPLICATION NUMBER: 60/312,858
PRIOR APPLICATION NUMBER: 60/271,840
PRIOR FILING DATE: 2001-08-16
PRIOR APPLICATION NUMBER: 60/271,840
PRIOR FILING DATE: 2001-02-27
PRIOR FILING DATE: 2001-02-27
                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
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SOFTWARE: PatentIn Ver. 2
SEQ ID NO 107
LENGTH: 7097
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PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: 60/286,096
PRIOR FILING DATE: 2001-04-21
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PRIOR FILING DATE: 2001-08-29
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PRIOR FILING DATE: 2001-06-20
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ORGANISM: Homo sapiens
  3676
                                                                                                                                                                                              3556 CCTAGCCCTCGGCAGGCCAGGCGCGAGCCCAGTTTACATCAAGTGGTGCTACAGCCC 3615
                                                                                            3616 TCCCGGCTCTCACCTCTGACCCAAAGCCCCCTCAGCTCCCGCACCGGCTCCCCTGAGCTC 3675
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                                              352 GCTCCGCACCACCATCCT 369
                                                                                                                                                                                                                     232 CCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCC 291
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GCCGCCCGTGCCCGGCCT 3693
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                                                                                                                                                                                                                                                                                                                                         Length 7097;
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RESULT 4 US-10-266-829-18

Sequence 18, Applica GENERAL INFORMATION:

Application US/10266829

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                                                                                                                                                                                                             US-10-281-024-16
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CURRENT APPLICATION NUMBER: US/10/266,829
CURRENT FILING DATE: 2002-10-09
PRIOR APPLICATION NUMBER: 09/756,168
PRIOR FILING DATE: 2001-01-09
PRIOR FILING DATE: 2000-07-20
PRIOR FILING DATE: 2000-07-20
PRIOR APPLICATION NUMBER: 60/145,220
PRIOR APPLICATION NUMBER: 60/145,220
PRIOR FILING DATE: 1999-07-23
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Best Local S
Matches 56
                                                                                                                                                                                                                         SEQ ID NO 16
LENGTH: 1372
TYPE: DNA
ORGANISM: Oryza sativa L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 16, Application US/10281024 GENERAL INFORMATION:
                                                                                                                                  Matches
                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: NICOTIANAMINE SYNTHASE AND GENE ENCODING TITLE OF INVENTION: THE SAME FILE REFERENCE: 55107 (71526) CURRENT APPLICATION NUMBER: US/10/281,024 CURRENT FILING DATE: 2002-10-25 PRIOR APPLICATION NUMBER: US/09/674,337A PRIOR APPLICATION NUMBER: US/09/674,337A PRIOR APPLICATION NUMBER: PCT/JP99/02305
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SOFTWARE: PatentIn Ver. 2
SEQ ID NO 18
                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Satoshi MORI
APPLICANT: Kyoko Higuchi
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TITLE OF INVENTION: 29 Human secreted proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: SITE
LOCATION: (1855)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 1860
158 GAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCAGCGGGCC
                                                    835 CCCGAGGACGTCAGGCGTGGCGGGTTCGACGTTCTGGCGGTGTGCCACCCGGAGGACGAG 894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          455 TCTTGGGGGCTTGTTGATTGGTGCTGGCTGTGCCCTCTACCCCTTGGGCTGGGACAGTGA 51
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                                                                                       Local Similarity
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                                                                                                                                                 Score 29.4; D
Pred. No. 2.9;
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Pred. No. 1.2;
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; ORGANISM: Homo Sapien
US-10-131-813A-293
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US-10-131-813A-293/c
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PRIOR APPLICATION NUMBER: 60/
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                               281 TGCAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCC
667 CGGGCCCCCTCTGCTTCTGAGAATGCGATGTTCAGGGCATACTGGATGGCCAGCCCAGTC
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CURRENT APPLICATION NUMBER: US/10/131,823A
CURRENT FILLING DATE: 2002-04-24
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PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
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PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-08-26
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PRIOR FILING DATE: 1997-09-17
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PRIOR APPLICATION NUMBER: 60/059113
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ORGANISM: Homo
281 TGCAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCC
                                                                                                        221 TCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGC
                                                                                                                                                  787 CCAATGGCAAAGATTAGGATGCCCGTGTCCCGTGCCTTAGCAGCCACCTCGGCCACGGAG
                                                                                                                                                                                           161 CAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCC 220
                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/059352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 1997-09-18
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                                                                 TCCTGAGGTCTCCCATCTGTCACGATCATTATGACCCGTGGCACATTCTCCCTCAGGGGC
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Godowski, Paul J.
Gurney, Austin L.
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                                                                                                                                                                                                                                                      7.8%;
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                                                                                                                                                                                                                                   Score 28.8; DB 6; Length 3 Pred. No. 7.5; Indels 0; Mismatches 113; Indels
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; ORGANISM: HOMO
US-10-131-824A-293
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                                                                                                                                           Query Match
Best Local :
                                                                                                                                                                                                                                                    SEQ ID NO 293
LENGTH: 3449
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GENERAL INFORMATION
                                                                                                                              Matches
                                                                                                                                                                                                                                                                                    Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/0
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/0
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: P3330R1C126
CURRENT APPLICATION NUMBER: US/10/131,824A
CURRENT FILING DATE: 2002-04-24
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 1997-09-19
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221 TCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTTGCATCAGCTGC 280
                                                                        161 CAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCC 220
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FILING DATE: 1997-09-19
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FILING DATE: 1997-09-17
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                                                     CCAATGGCAAAGATTAGGATGCCCGTGTCCCGTGCCTTAGCAGCCACCTCGGCCACGGAG 728
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                                                                                                                                              Similarity
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Wood, William
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Godowski, Paul J.
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                                                                                                                            Conservative
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Pred. No. 7.
                                                                                                                        Mismatches 113;
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                                                                                                                                          ; ORGANISM: Homo Sapier US-10-131-826A-293
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US-10-131-826A-293/c
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LENGTH: 3449
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Best Local
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PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
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                    161 CAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCC 220
787 CCAATGGCAAAGATTAGGATGCCCGTGTCCCGTGCCTTAGCAGCCACCTCGGCCACGGAG 728
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FILING DATE: 1997-09-17
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Wood, William
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Godowski, Paul J.
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Filvaroff, Ellen
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                                                                       Conservative
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                                                                   Score 28.8; DE Pred. No. 7.5; 0; Mismatches
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US-10-131-829A-293
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US-10-131-829A-293/c
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LENGTH: 3449
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                                                                                           Query Match
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PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C138
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Remaining Prior Application d
NUMBER OF SEQ ID NOS: 550
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                  161 CAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCC 220
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                                                                        Local
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APPLICATION NUMBER: 60/059122
FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/059184 FILING DATE: 1997-09-17
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Godowski, Paul J.
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                                                       Conservative
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                                                                      7.8%;
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                                                      Score 28.8; DF Pred. No. 7.5; O; Mismatches
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US-10-125-926A-293/C
                                                                                US-10-125-926A-293
                                                                                                                                    Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 550 SEQ ID NO 293 LENGTH: 3449
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GENERAL INFORMATION
  Matches
                  Query Match
Best Local Similarity
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263
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PRIOR FILING DATE: 1997-09-17
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PRIOR APPLICATION NUMBER: 60/059117
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                                                                                           TYPE: DNA
ORGANISM: Homo Sapien
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                                                                                                                                                                                                                                                                                                    FILING DATE: 1997-09-
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LENGTH: 3449
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PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C85
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                                                                                                TYPE: DNA
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                                                                                                                                                                                       FILING DATE: 1997-09-19
APPLICATION NUMBER: 60/059588
FILING DATE: 1997-09-19
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FILING DATE: 1997-09-18
APPLICATION NUMBER: 60/059352
                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/059184 FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/059122
FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/059117 FILING DATE: 1997-09-17
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Sherwood, Steven
Smith, Victoria
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Godowski, Paul J.
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                                                                           Sapien
   7.8%;
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 Pred.
                   Score
28.8;
No. 7
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                 6;
               Length 3449;
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                                                                   Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 550 SEQ ID NO 293
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/127,831A CURRENT FILING DATE: 2002-10-15 PRIOR APPLICATION NUMBER: 60/049911 PRIOR FILING DATE: 1997-06-18
                                                                                                                              PRIOR APPLICATION NUMBER: 60/059588 PRIOR FILING DATE: 1997-09-19
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PRIOR
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               ORGANISM: Homo Sapien
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                                                   LENGTH:
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                                                                                                                                                                                                      FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/
FILING DATE: 1997-09-18
                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/059122
FILING DATE: 1997-09-17
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FILING DATE: 1997-09-17
                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/059184
                                                                                                                                                                                    APPLICATION NUMBER: 60/059352
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Sherwood, Steven
Smith, Victoria
Stewart, Timothy A.
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Wood, William
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Godowski, Paul J.
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Filvaroff, Ellen
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US-10-127-835A-293/c
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GENERAL INFORMATION:
APPLICANT: Baker Kevin P.
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Best Local (
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Remaining Prior Application data
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 293
LENGTH: 3449
TYPE: DNA
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APPLICANT:
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CURRENT APPLICATION NUMBER: US/10/127,835A
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
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                                                                                                             PRIOR FILING DATE: 1997-09-19
                                                                                                                                  PRIOR APPLICATION NUMBER: 60/059588
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Local Similarity 45.9%;
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Watanabe, Colin K
Wood, William
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Godowski, Paul J.
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0; Mismatches 113; Indels
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                                                                                        removed - See File Wrapper or PALM
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ORGANISM: Homo Sapien

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                                                                CGGGCCCCCTCTGCTTCTGAGAATGCGATGTTCAGGGCATACTGGATGGCCAGCCCAGTC
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Search completed: November 7, 2002, 17:56:44 Job time: 28.3091 secs

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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## REFERENCE AUTHORS TITLE JOURNAL KEYWORDS SOURCE ACCESSION VERSION LOCUS DEFINITION RESULT 1 AC079773/c Result No. O a O ი 000 000 00000 G ORGANISM Homo sapiens AC079773 AC079773.8 G Eukaryota; Metazoa; Chordata; Craniata; Mammalia; Eutheria; Primates; Catarrhin 1 (bases 1 to 129676) Sulston, J.E. and Waterston, R. Toward a complete human genome sequence Genome Res. 8 (11), 1097-1108 (1998) Homo sapiens. Homo sapiens HTG Query Match Length 97.0 3.3 444 3.8 70395 3.7 94718 3.7 167693 3.7 176967 3.7 180837 3.7 180837 3.7 192464 3.7 200145 5.5 256 98506 98506 111900 111914 1114382 1154363 126363 126363 126363 158728 163932 163932 194402 194402 198863 206648 GI:15145561 3190 43570 86093 90020 94029 1388 1982 2666 129676 bp DNA linear PRI 09-JA BAC clone RPI1-258B17 from 2, complete sequence. ВВ AC126575 AC098517\_0 AL596180 AC103498 AC094168 AC0123419 ATHAP2A AY065052 AY065057 AB035577 AB019231 AC114102 ATT24H18 AC117309 ATT2L20 AC096152 AC107560 AC102588 AC123116 AC094585 AC022824 AC009679 AC090971 AC126666 AX393645 AC011639 AC022742 AC112072 AC099984 AC026770 AX136281 AX358762 AX362255 AX136698 AX319942 AX150120 AF034633 ALIGNMENTS SUMMARIES Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo. AX319944 Sequence AX150120 Sequence AX150120 Sequence AX150120 Sequence AX150120 Sequence AX160120 Rattus no AC099984 Mus muscu AC026770 Homo sapi AC01639 Homo sapi AC022824 Homo sapi AC029971 Homo sapi AC09971 Homo sapi AC126666 Rattus no AX393645 Sequence Y13720 Arabidops AY065052 Arabidops AY065052 Arabidops AY065052 Arabidops AX194474 Nicotiana AF035577 Amblyomen AB019231 Arabidops AC117309 Rattus no AL353013 Arabidops AC117309 Rattus no AL3596180 Mouse DNA AC103498 Rattus no AC094168 Rattus no AC094168 Rattus no AC0123419 Rattus no AC0123419 Rattus no AC123419 Rattus no AC124653 Homo sapi AC012568 Mattus no AC012568 Rattus no AC012568 Rattus no AC012568 Rattus no AC012568 Rattus no AC012568 Rattus no AC012568 Rattus no AC10316 Rattus no AC10316 Rattus no AC10316 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC103168 Rattus no AC079773 Description PRI 09-JAN-2002 Homo sapi Homo sapi Sequence Sequence Sequence Sequence Sequence

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The sequence of Homo sapiens BAC clone RP11-258B17
Unpublished (2001)
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Submitted (09-AUG-2001) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St.
MO 63108, USA
5 (bases 1 to 129676)
                                                                                                                                                                                                                                              The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Frengen, E., Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at the Roswell Park Cancer Institute (http://bacpac.med.buffalo.edu)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (09-JAN-2002) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 63108, On Aug 9, 2001 this sequence version replaced gi:14488388.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct Submission
Submitted (10-SEP-2000) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St. Louis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.
                                                                                                                                            NEIGHBORING SEQUENCE INFORMATION:
The clone sequenced to the left is RP11-1L22; the clone sequenced to the right is RP11-159N20, 2000 bp overlap. Actual start of the clone is at base position 1 of RP11-258B17; actual end is at base
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. Louis MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc
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                                                          RP11-258B17 contains a single plasmid region Location/Qualifiers
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                                                                                                                           position 60003 of RP11-159N20.
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Contact: sapiens@watson.wustl.edu
----- Summary Statistics
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/organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                 Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Series: IRAL Plate: 25 Row: n Column: 18
This clone was selected for full length sequencing because it passed the following selection criteria: Hexamer frequency ORF
                                                                                                                                                                                                                                                                                                                                                                                                                                          CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LINL)
DNA Sequencing by: Institute for Systems Biology
http://www.systemsbiology.org
contact: amadandsystemsbiology.org
Anup Madan, Rachel Dlckhoff, Jessica Fahey, Stephanie Ford, Julia
Greene, Mark Ketteman and Anuradha Madan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Direct Submission
Submitted (25-JUL-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 826)
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/tissue_type="Skin, melanotic me
/clone_lib="NIH_MGC_20"
/lab_host="DH10B-R"
/note="Vector: pOTB7"
a 197 c 228 g 164 t
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Direct Submission
Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team);
Kazusa-Kamatari, Kisarazu, Chiba 292-0812, Japan
(E-mail:genomics@hri.co.jp, Tel:81-438-52-3975, Fax:81-438-52
NEDO human cDNA sequencing project supported by Ministry of
                                                                                                                                                                                                                                                                     oligo capping; fis (full insert sequence). Homo sapiens alzheimer cortex cDNA to mRNA,
                                                                                                                                                                                                                                                                                                                                                  Homo
                                                                                  Unpublished
2 (bases 1
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Isogai,T. and Yamamoto,J.
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                 Fax:81-438-52-3986)
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Location/Qualifiers
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NCTVNVQDMCQKEVMEQSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRPKKRGSSASALRPGLRTTILFLKLALFSAHC"
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215. .640
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/db_xref="taxon:9606"
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/db_xref="GI:21753576"
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/clone_lib="BRALZ2"
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                                                    AATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACC
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                                        AATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACC
                                                                                          GACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAACACAGCATGAGAACACAGTAAATA
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Patent: EP 1067182-A 203 10-JAN-2001;
Helix Research Institute (JP)
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1890)

Ota, T., Isogai, T., Nishikawa, T., Kawai, Y., Sugiyama, T. and
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/db_xref="taxon:9606"
274. .771
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Sequence
AX358762
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Secreted and transmembrane polypeptides and nucleic a
            AX362255.1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
433 c 435 g 33
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AX136698
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Homo sapiens
                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; (bases 1 to 587)
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                                                                                         Ota,T., Isogai,T., Nishikawa,T., Kawai,Y.,
Hayashi,K.
                                                              Secretory protein or membrane protein Patent: EP 1067182-A 620 10-JAN-2001;
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/db_xref="taxon:9606"
433 c 435 q 33'
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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AX319942.1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                         /translation="MASPSLPGSDCSQIIDHSHVPEFEVATWIKITLILVYLIIFVMG
LLGNSATIRVTQVLQKKGYLQKEVTDHMVSLACSDILVFLIGMETYSIIWHELTTS
SYTLSCKLHTFLEEACSYATLLHVLTLSFERYIAICHPFRYKAVSGPCQVKLLIGFVW
VTSALVALPLLFAMGTEYPLVNVPSHRGLTCNKSSTRHHEQPETSNMSICTNLSSRWT
VTSALVALPLLFLAMGTEYPLVNVPSHRGLTCNKSSTRHHEQPETSNMSICTNLSSRWT
VTSSLIGGAFVYYLVVLSVARWGWNMQVLMKSQKGSLAGGTRPPQLRKSESEESRT
ARRQTIIFLRLIVYTLAVCWMPNQIRRIMAAAKPKHDWTRSYFRAYMILLPFSETFFY
ASRQTIIFLLYTYSSQQERRYFYQVLCCRLSLQHANHEKRLRVHAHSTTDSARRYQRPL
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                                                                                                                                                                                                        /codon_start=]
                                                                                                                                                                                                                     /note="unnamed
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from Patent WO0181634.
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Submitted
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MCKee, K.K., Tan, C.P., Palyha, O.C., Liu, J., Feighner, S.D.,

Hreniuk, D.L., Smith, R.G., Howard, A.D. and Van der Ploeg, L.H.

Cloning and characterization of two human G protein coupled

receptor genes (GPR38 and GPR39) related to the growth hormo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      complete
AF034633
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McKee, K.K., Tan, C.P.,
                                                                                                                                                                                                                                                                                                                                           secretagogue and neurotensin receptors
Genomics 46 (3), 426-434 (1997)
98110578
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                      d (17-NOV-1997) Biochemistry and Physiology, Merck and
Box 2000, Rahway, NJ 07065, USA
Location/Qualifiers
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                          hormone secretagogue
                                                                                                              /map="2q21-22"
/tissue_type="brain"
/dev_stage="fetal"
/product="GPR39"
                                        /note-"orphan G
                                                        /gene="GPR39"
                                                                                   /gene="GPR39"
                                                                                                                                                         /chromosome="2"
                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
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Methods and compositions for the diagnosis and
Methods and compositions for the diagnosis and
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Sequence 3 from Patent W00181634.
                                                                                                                                                                                                    Millennium Pharmaceuticals,
Location/Qualifiers
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Patent: WO 0181634-A 3 01-NOV-2001
                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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/db_xref="taxon:9606"
42. .1403
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VFQSSLIFGAFVVLVVLLSVAFMCWNMMQVLMKSGKGSLAGGTBAPQLLKSESSESRT
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ARRQTIFFLKLIVYTLAVCKWHOMIRRIMAAKKHHDWTRYFRAYMILLFESTFFY
ARRQTIFFLKLIVYTLAVCKWHOMIRRIMAAKKHHDWTRYFRAYMILLFESTFFY
ARRQTIFFLKLIVYTLAVGWHOMIRRIMAAKKHRYHAHSTTDSARFYQRPL
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SYTLSCKLHTFLFEACSYATLLHVLTLSFERYIAICHPFRYKAVSGPCQVKLLIGFVM
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                                                                                                                                                                                                                                                                                                                                                                                                                                GI:17901491
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Pred. No. 1.2e-207;
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VTSALVALPLLFAMGTEYPLVNVPSHRGLTCNRSSTRHHEQPETSNMSICTNLSSRWI

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                        241 GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGGCGCAGC 300
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1 (bases 1 to 444)

Kroes, R.A., Moskal, J.R. and Yamamoto, H.

Differential gene expression in cancer Patent: WO 0136685-A 95 25-MAY-2001; NYXIS NeuroTherapies, Inc. (US)
                                                                                                                                                                                                                                                                                                                                                                                            AX150120.1 GI:14348148
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/db_xref="taxon:9606"
113 c 94 g 14:
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ARRQTIIFLRLIVVTLAVCWHOQIRR MAAAKFKHDWTRSYFAAYMILLPFSETFFY
LSSVLMPLLYTVSSQQFRRVFYQVLCCRLSLQHANHEKRLRVHAHSTTDSARFYVQPL
LFASRRQSSARRTEKIFLSTFQSEAEPQSKSQSLSLESLEPNSGAKPANSAAENGFQE
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Pred. No. 2e-203;
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RESULT 12 ACCLISO72 LOCUS DEFINITION ACCESSION VERSION KEYMORDS SOURCE ORGANISM REFERENCE AUTHORS	7 0	Qy .504 Db 131	Qy 444 Db 191	Qy 384 Db 251	Db 311
AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC112072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072  AC12072	GTTGGCTGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG 29	TTGACAGCTTCTTTTGCCACAAGCAAGAGAATTTAACACTCT  - - - - - - - - - - - - - - - - - - -	TITAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTGGA 503	GAGACAGAGAAAAAAACACAGCATGAGAACACAGTAAATGAATAAAACCATAAAATA 443 	

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (17-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 13, 2002 this sequence version replaced gi:20303189.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (19-FEB-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 92874)
Worley, K.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Worley, K.C.
Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NOTE: Estimated insert size may differ from sequence length (see http://www.bysc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 50 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 as soon as it
be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Center: Baylor College of Medicine Center code: BCM
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27; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                            Submitted (22-NOV-2001) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA All repeats were identified using RepeatMasker: Smit, A.F.A. & Green, P. (1996-1997)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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           NOTE: This record contains 84 individual sequencing reads that have not been assembled into contigs. Runs of N are used to separate the reads and the order in which they appear is completely arbitrary. Low-pass sequence sampling is useful for identifying clones that may be gene-rich and allows overlap relationships among clones to be deduced.
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                                                                                                                                                                                         Contact: sequence_submissions@genome.wi.mit.edu
------ Project Information
Center project name: L13721
Center clone name: 24_D_9
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55385: contig of 7
6 55485: gap of 10
6 56162: contig of 6
3 56262: gap of 10
                                                                                                                                                                                                                                                                                                                                                        52194: gap of 100 bp
52922: contig of 728 bp in length
53022: gap of 100 bp
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30249: gap of 100 k
30990: contig of 741
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100 bp
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51307: gap of 699 bp 1
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100 bp
17101: contig of 739 bp 1
17101: gap of 100 bp
57848: contig of 747 hr
1948: gap of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     42052: gap of 100 bp 101 length 42893: contig of 741 bp in length 42893: contig of 730 bp in length 43623: contig of 751 bp in length 4474: contig of 751 bp in length 4574: gap of 100 bp 4574: gap of 100 bp in length 45323: contig of 749 bp in length 4523: contig of 73 bp in length 46156: contig of 73 bp in length 46256: gap of 100 bp 46256: gap of 100 bp in length 46256: gap of 100 bp in length 47152: gap of 100 bp in length 47152: gap of 100 bp in length
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48723: contig of 737 bp
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49570: contig of 747 bp
9670: gap of 100 bp
50408: contig of 738 bp
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ap of
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contig of 715 bp
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of 734 bp in length
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AC026770/c
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383 AGAGACAGAGAAAAGAAAAACACA 406
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Submitted (24-MAR-2000) Multimegabase Sequencing Center, University of Washington, PO Box 357730, Seattle, WA 98195, USA
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On NOV 17, 2000 this sequence version replaced g1:10697426
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (17-NOV-2000) Multimegabase Sequencing Center, Institute for Systems Biology, 4225 Roosevelt Way NE, Suite 200, Seattle, WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rowen,L., Madan,A., Qin,S., Baradarani,L., Birditt,B., Bloom,S., Dors,M., Dickhoff,R., Fleetwood,P., Harrison,G., Kaur,A., Madan,A., Nesbitt,R., Traicoff,R. and Hood,L.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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94718 bp DNA linear PRI 17-NOV-20
Homo sapiens chromosome 15 clone CTD-2308G16 map 15q21, complete
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Assembly program: Phrap;
                                                                                                                                          /note="Overlap with RP11-313P18, 66240. .72320
                                                                                                                                                                                                                                                                                                                              /Clone_lib="Cal Tech human BAC library 11"
/note="This clone overlaps RP11-313P18, RP11-414J1 and RP11-56B16. Data from all of these BACs were combined,
                               /note="number of Ts unclear"
72300 . .72320
/note="low quality data"
72960 . .72980
                                                                                                                                                                                                                                                                                  CTD-2306G16.
                                                                                                                                                                                                                                                                                           with an attempt to generate the consensus sequence from CTD-2306G16."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
1. .94718
note="low quality data"
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                                                                                                                                                                                                                        note="overlap with RP11-414J1, AC020685"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (08-OCT-1999) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On May 4, 2001 this sequence version replaced gi:13123327. All repeats were identified using RepeatMasker:
Smit, A.F.A. & Green, P. (1996-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            http://ftp.genome.washington.edu/RM/RepeatMasker.html
------ Genome Center
Center: Whitehead Institute/ MIT Center for Genome Research
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                  Sequencing vector: M13; M77815; 12% of reads
Sequencing vector: M13; M77815; 12% of reads
Sequencing vector: Plasmid; n/a; 88% of reads
Sequencing vector: Plasmid; p/a; 80% of reads
Sequencing vector: Plasmid; n/a; 80% of reads
Sembly program: Phrap; version 0.960731
Consensus quality: 165314 bases at least Q30
Consensus quality: 165712 bases at least Q30
Consensus quality: 165713 bases at least Q30
Insert size: 170000; agarose-fp
Insert size: 166933; sum-of-contigs
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                                                                                                                                                                                                                                                                                                                                                                            Contact: sequence_submissions@genome.wi.mit.edu
                                                                                                                                                                                                                                                                                                                   Center project name: L3327
Center clone name: 13_H_19
                                                                                                                                                                                                                                                                                                                                                                                                                                   Web site: http://www-seq.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Center code: WIBR
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/note="low quality data"
a 19737 c 19836 g 26384
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88020. .88070
coverage:
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100.0%; Pred. No. 0.
tive 0; Mismatches
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7.8 in
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Q20 bases; agarose-fp
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WORKING DRAFT SEQUENCE,
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RESULT 16
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                    ORGANISM
                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local
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                                                                                                                                                                                                                                                                                               382 GAGAGACAGAGAAAAGAAAAACA 404
                                                                                                                                    ACO22742 176967 bp DNA linea
Homo sapiens chromosome 15 clone RP11-762A1 map
                                                                                               AC022742
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
                    Homo sapiens
                                                           HTG; HTGS_PHASE1;
                                                                           AC022742.4 GI:13123383
                                                                                                                    SEQUENCE, 4 unordered pieces.
                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                    Similarity
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NOTE: This is a 'working draft' sequence. It currently consists of 8 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         be preserved
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90240 120347: contig of 30108 bp in length
120348 120447: gap of 100 bp
120448 167227: contig of 46780 bp in length
167228 167327: gap of 100 bp
167328 167693: contig of 366 bp in length.
Location/Qualifiers
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54882 54981: gap of 100 bp
54982 57393: contig of 2412 bp in length
57394 57493: gap of 100 bp
57494 64870: contig of 7377 bp in length
64871 64970: gap of 100 bp
64871 73018: contig of 8048 bp in length
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                                                                                                                                                                                                                                                                                                                                     Conservative
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90240. .120347
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="RP11-13H19"
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167328, .167693
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1. .54881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="assembly_fragment
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                                                                                                                                                                                                                                                                                                                                                      3.7%; Score 23; 100.0%; Pred. No.
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                                                           HTGS_DRAFT; HTGS_FULLTOP
                                                                                                                                                                                                                                                                                                                                   0,
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0.12;
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                                                                                                                                                            linear
                                                                                                                                        15,
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WORKING DRAFT
    Euteleostomi;
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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequencing vector: M13; M77815; 2% of reads
Sequencing vector: Plasmid; n/a; 98% of reads
Sequencing vector: Plasmid; n/a; 98% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.960731
Consensus quality: 176032 bases at least Q40
Consensus quality: 176015 bases at least Q20
Consensus quality: 176615 bases at least Q20
Insert size: 186000; agarose-fp
Insert size: 176667; sum-of-contigs
Quality coverage: 9.0 in Q20 bases; agarose-fp
Quality coverage: 9.5 in Q20 base
* NOTE: This is a 'working draft' sequence. It currently
consists of 4 contigs. The true order of the pieces
* is not known and their order in this sequence record is
* arbitrary. Caps between the contigs are unknown.
* This record will be updated with the filshed sequence
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Submitted (06-FEB-2000) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On Feb 25, 2001 this sequence version replaced gi:11990718.
All repeats were identified using RepeatMasker:
Smit, A.F.A. & Green, P. (1996-1997)
http://ftp.genome.washington.edu/RM/RepeatMasker.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zimmer, A. and Zody, M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   as soon as it is available and the accession number will be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: sequence_submissions@genome.wi.mit.edu
----- Project_Information
Center project name: L5677
Center clone name: 762_A_1
                                                                                                                                                                                                                                                                                               14401 14500: gap of 100 bp 14501 61165: contig of 46665 bp in length 61166 61265: gap of 100 bp 61266 112521: contig of 51256 bp in length 112522 112621: gap of 100 bp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Web site: http://www-seq.wi.mit.edu
                /clone_lib="RPCI-11 Human Male BAC"
                                                 /clone="RP11-762A1"
                                                                              /chromosome="15"
/map="15"
                                                                                                                                                                      ∕organism="Homo sapiens"
                                                                                                                                         /db_xref="taxon:9606"
                                                                                                                                                                                                                                    ocation/Qualifiers
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AUTHORS
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KEYWORDS
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Naylor, J., Nguyen, C.,
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Db 166420 GAGAGACAGAGAAAAGAAAAACA 166398
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                                                                                                                                                                                                                                                                                                                 Submitted (06-FEB-2000) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA 3 (bases 1 to 180837)
Birren,B., Nusbaum,C., Lander,E., Ali,A., Allen,N., Anderson,S., Barna,N., Bastien,V., Bloom,T., Boguslavkiy,L., Boukhgalter,B., Camarata,J., Chang,J., Chazaro,B., Choepel,Y., Collymore,A., Cook,A., Cooke,P., DeArellano,K., Dear,K., Diaz,J.S., Dodge,S.,
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Homo sapiens chromosome 15 clone
SEQUENCE, 8 unordered pieces.
ACQ22824
Faro, S., Ferreira, P., FitzGerald, M., Gage, D., Galagan, J., Gardyna, S., Gord, S., Graham, L., Grand-Pierre, N., Hagos, B., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, Karatas, A., Kells, C., Landers, T., Levine, R., Lindblad-Toh, K. Liu, G., MacLean, C., Macdonald, P., Major, J., Matthews, C., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., Matthews, C., Magella, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matth
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1 (bases 1 to 180837)
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112622. .176967
/note="assembly_fragment"
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61266. .112521
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100.0%; Pred. No.
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                                                                                                                 Lindblad-Toh, K.,
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WORKING DRAFT
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Nicol, R.,

Norman, C.H.,

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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            O'Connor,T., O'Donnell,P., O'Neil,D., Oliver,J., Peterson,K., Phunkhang,P., Pierre,N., Raymond,C., Retta,R., Rise,C., Rogov,P., Roman,J., Roy,A., Schauer,S., Schupback,R., Seaman,S., Severy,P., Smith,C., Spencer,B., Stange-Thomann,N., Stojanovic,N., Talamas,J., Tesfaye,S., Theodore,J., Topham,K., Travers,M., Vassiliev,H., Tesfaye,S., Theodore,J., Topham,K., Travers,M., Vassiliev,H., Vo,A., Wilson,B., Wu,X., Wyman,D., Young,G., Zainoun,J.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Center project name: L6128
Center clone name: 81_1_17
Center clone name: 81_1_17
Sequencing vector: M13: M77815; 3% of reads
Sequencing vector: M13: M77815; 3% of reads
Sequencing vector: M13: M77815; 3% of reads
Sequencing vector: M13: M77815; 0% of reads
Chemistry: Dye-terminator Big Dye; 110% of reads
Assembly program: Phrap; version 0.960731
Consensus quality: 177875 bases at least Q40
Consensus quality: 179781 bases at least Q20
Consensus quality: 179781 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Insert size: 182000; agarose-fp
Insert size: 180137; sum-of-contigs
Ouality coverage: 7.8 in 020 bases; agarose-fp
Quality coverage: 7.8 in 020 bas
NOTE: This is a 'working draft' sequence. It currently
consists of 8 contigs. The true order of the pieces
is not known and their order in this sequence record is
arbitrary. Gaps between the contigs are represented as
runs of N, but the exact sizes of the gaps are unknown.
This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: sequence_submissions@genome.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Center code: WIBR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Center: Whitehead Institute/ MIT Center for Genome Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Web site: http://www-seq.wi.mit.edu
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                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                     /clone_lib="RPCI-11 Human Male BAC"
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                                                                                                                                                                                                                                                                                      lone="RP11-81I17"
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COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Birren, B., Linton, L., Nusbaum, C., Lander, E., Allen, N., Anderson, M. Baker, J., Baldwin, J., Barna, N., Beckerly, R., Benn, J., Brown, A., Castle, A., Cerny, J., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Depayre, E., Devon, K., Dewar, K., Donelan, L., Doyle, M., Ferreira, P., FitzHugh, W., Forrest, C., Funke, R., Gage, D., Galagan, J., Gardyna, S., Gilbert, D., Grant, G., Hagos, B., Heaford, A., Horton, L., Howland, J.C., Jones, C., Kann, L., Karatas, A., Lehoczky, J., Lieu, C., Locke, K., Macdonald, P., Karatas, A., Lehoczky, J., Lieu, C., Locke, K., Macdonald, P., Marquis, N., McEwan, P., McGurk, A., McKernan, K., McLaughlin, J., Meldrim, J., Mollan, M., Morris, W., Morrow, J., Mychalecky J., Naylor, J., Niloff, M., O'Connor, T., O'Donnell, P., Pavlin, B., Naylor, J., Niloff, M., O'Connor, T., Roberts, D., Roy, A., Severy, P., Characom, K., Pollara, V., Riley, R., Roberts, D., Roy, A., Severy, P., Characom, A., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom, C., Characom,
                                                                                                                                                                                                                                                                                                                                                                                      submitted (28-AUG-1999) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On Apr 20, 2000 this sequence version replaced gi:6067177. All repeats were identified using RepeatMasker: Smit, A.F.A. & Green, P. (1996-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stange-Thomańn,N., Stojanovic,N., Stone,C., Subramanian,A., Tesfaye,S., Torruella-Miller,I., Vassiliev,H., Vo,A., Wagner,A., Wheeler,J., Wu,X., Wyman,D., Ye,W.J. and Zody,M.
                                                                                                                                                                                                                                                                                                                       http://ftp.genome.washington.edu/RM/RepeatMasker.html
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AC009679.3 GI:7622340
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Direct Submission
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Unpublished
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                                                                  Center project name: L2273
Center clone name: 56_B_16
                                                                                                                                         ------ Project Information
                                                                                                                                                                      Contact: sequence_submissions@genome.wi.mit.edu
Sequencing vector: M13; M77815;
                                                                                                                                                                                                               Web site: http://www-seq.wi.mit.edu
                                                                                                                                                                                                                                                Center code: WIBR
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40288. .68512
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21160. .40187
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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100.0%; Pred. No.
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WORKING DRAFT SEQUENCE, 27 unordered
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## 156 9617: contig of 1462 bp in le 9618 9717: gap of 100 bp 9718 11851: contig of 2134 bp in ler. 11852 11951: gap of 100 bp 11952 14299: contig of 2348 bp in ler. 14300 14399: gap of 100 bp 14400 16295: contig of 3448 bp in len. 16296 16305: contig of 100 bp 16296 16305: contig of 100 bp 16296 16305: contig of 100 bp 16396; gap of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp 16396; contig of 100 bp
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                                                                                    63598 74774

74775 74844: 9

74775 74874: 9

74775 74874: 9

89189 89288: 9

89289 107293: 107294 107293: 107294 107293: 12450: 12450: 12450: 12450: 124531: 141531
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Quality coverage: 3.4 in Q20 bases; agarose-fp Quality coverage: 3.5 in Q20 bases; sum-of-contigs
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Insert size: 181885; sum-of-contigs
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                                                                                                                                                                                                                                                                                        55755 55854: gap of 100 bg
55855 63497: contig of 7643
63498 63597: gap of 100 bg
63598 74774: contig of 11177
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18 18517: gap of 100 bp
18 20760: contig of 2243 bp
51 20860: gap of 100 bp
51 20860: gap of 3016 bp
77 23976: gap of 100 bp
77 23976: contig of 3636 bp
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7 184485: contig of Location/Qualifiers
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8055: contin ~ 6
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40530, .47290
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163367. .184485
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36984 c 37799 g 54422
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23977. .27612
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20861. .23876
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18518. .20760
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141632. .163266
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31061. .34289
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107394. .124850
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On Nov 27, 2001 this sequence version replaced gi:13624393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pate,D. and Hood,L.
Direct Submission
Submitted (27-NOV-2001) Multimegabase Sequencing Center, Institute for Systems Biology, 4225 Roosevelt Way NE, Suite 200, Seattle, WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rowen, L., Madan, A., Qin, S., Baradarani, L., Birditt, B., Bloom, S. Burke, J., Dors, M., Fleetwood, P., Kaur, A., Madan, A., Nesbitt, R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (21-MAR-2001) Multimegabase Sequencing Center, Institute for Systems Biology, 4225 Roosevelt Way NE, Suite 200, Seattle, WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rowen,L., Madan,A., Qin,S., Baradarani,L., Birditt,B., Bloom,S., Dors,M., Dickhoff,R., Fleetwood,P., Harrison,G., Kaur,A., Madan,A., Nesbitt,R., Traicoff,R. and Hood,L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequencing of human chromosome 15 D15S146-D15S117 region unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rowen,L., Madan,A., Qin,S., Baradarani,L., Birditt,B., Bloom,S., Burke,J., Dors,M., Fleetwood,P., Kaur,A., Madan,A., Nesbitt,R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 192464)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pate, D. and Hood, L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AC090971.3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (bases 1 to 192464)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chemistry: Dye-terminator Big Dye; 90% of reads Chemistry: Dye-primer Big Dye; 10% of reads Assembly program: Phrap; version 0.990399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Center: Multimegabase Sequencing Center
Center code: UWMSC
Web site: http://chroma.mbt.washington.edu/msg_www
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: leerowen@systemsbiology.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequencing vector: pUC18; L08752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ----- Summary Statistics
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                                                                   /note="overlap with CTD-2650P22 AC090970" 80765. .192464
                                                                                                                                                                                                                                                                       /note-"Data from overlapping BACs CTD-2308G16, RP11-13H19, RP11-47K1, and CTD-2650P22 was added and the consensus sequence was determined from RP11-56B16 to the extent
                                                                                                                                                                                                                                                possible"
                                                                                                                                                                                                                                                                                                                                              /clone="RP11-56B16"
/clone_lib="RPCI human BAC library 11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                            note="overlap with RP11-47K1 AC016824"
                                                                                                                        /note="overlap with CTD-2308G16 AC026770"
76219. .192464
                                                                                                                                                                                              note="overlap with RP11-13H19 AC011639"
                                                                                                                                                                                                                                                                                                                                                                                                    /map="15q21.3"
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                                                                                                                                                                                                                             .167631
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                                                                                                                                                                                                                                                                                                                        Muzny, D.M., Adains, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbarla, J., Benton, J., Binage, K., Blankenburg, K., Bonnin, D., Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Burarla, J., Esenton, J., Erown, E., Brown, M., Bryant, N.P., Bunay, C., Burch, P., Brievett, C., Burch, J., Chavez, D., Chen, R., Chen, R., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, Z., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, Z., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, Z., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, Z., Chen, Z., Chen, Z., Chowdhry, I., Chert, R., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Chen, Z., Ch
                             Submitted (08-JUL-2002) Human Genome Sequencing Center, Depa of Molecular and Human Genetics, Baylor College of Medicine, Baylor Plaza, Houston, TX 77030, USA
                                                                                                                          Direct Submission
                                                                                                                                                       Worley, K.C.
                                                                                                                                                                                                                 Unpublished
                                                                                                                                                                                                                                                     Direct Submission
                                                                                                                                                                                                                                                                                  Wu,C., Wu,Y., Wu,Y.F., Zhou,J., Zorrilla,S., Nelson,D.,
Weinstock,G. and Gibbs,R.
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AC126666.1 GI:21703541
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Mammalia; Eutheria; Rodentia; Sclurognathi; Muridae; Murinae;
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(bases 1 to 200145)
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100.0%; Pred. No.

 Mismatches

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                                                                                          Department
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Worley, K.C.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NOTE: Estimated insert size may differ from sequence length
(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).
NOTE: This is a 'working draft' sequence. It currently
consists of 61 contigs. The true order of the pieces
is not known and their order in this sequence record is
arbitrary. Gaps between the contigs are represented as
runs of N, but the exact sizes of the gaps are unknown.
This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329
Consensus quality: 141149 bases at least Q40
Consensus quality: 153199 bases at least Q30
Consensus quality: 163387 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Center project name: GARB
Center clone name: CH230-3P12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              soon as it is available and the accession number will preserved.
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1377: gap of
2611: contig
2711: gap of
3823: contig
3923: contig
3927: contig
5557: gap of
5657: gap of
6844: qap of
9032: contig
9132: gap of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 34 from Patent WO0212329.
AX393645
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Unpublished
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31 c 52 g 8:
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                               /product="transcription factor"
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Cheuk,R., Chen,H., Kim,C.J., Koesema,E., Meyers,M.C., Banh,J.,
Bowser,L., Carninci,P., Chang,E., Dale,J.M., Goldsmith,A.D.,
Hayashizaki,Y., Ishida,J., Jones,T., Kamiya,A., Karlin-Neumann,G.,
Kawai,J., Lam,B., Lee,J.M., Lin,J., Miranda,M., Narusaka,M.,
Nguyen,M., Onodera,C.S., Palm,C.J., Quach,H.L., Sakurai,T.,
Satou,M., Seki,M., Southwick,A., Tang,C.C., Toriumi,M., Wu,H.C.,
Yamada,K., Yamamura,Y., Yu,G., Yu,S., Shinozaki,K., Davis,R.W.,
Theologis,A. and Ecker,J.R.
Arabidiopsis cDNA clones
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Cheuk, R., Chen, H., Kim, C.J., Koesema, E., Meyers, M.C., Banh, J.,
Cheuk, R., Chen, H., Kim, C.J., Koesema, E., Meyers, M.C., Banh, J.,
Bowser, L., Carninci, P., Chang, E., Dale, J.M., Goldsmith, A.D.,
Hayashizaki, Y., Ishida, J., Jones, T., Kamiya, A., Karlin-Neumann, G.,
Kawai, J., Lam, B., Lee, J.M., Lin, J., Miranda, M., Narusaka, M.,
Nguyen, M., Onodera, C.S., Palm, C.J., Quach, H.L., Sakurai, T.,
Sakiu, M., Southwick, A., Tang, C.C., Toriumi, M., Wu, H.C.,
Yamada, K., Yamamura, Y., Yu, G., Yu, S., Shinozaki, K., Davis, R.W.,
contributed equally contributed equally contributed equally contributed equally to this work as PIs.

Location/Qualifiers

1. 1982
                                                                                                                                                                                                                     The Salk, Stanford, PGEC (SSP) Consortium members carried out the sequencing and annotation of the RAFL cDNAs: Cheuk,R., Chen,H., Kim,C.J., Koesema,E., Meyers,M.C., Shinn,P., Banh,J. Bowser,L., Dale,J.M., Goldsmith,A.D., Jiang,P.X., Jones,T., Karlin-Neumann,G. Lam,B., Lee,J.M., Lin,J., Liu,S.X., Miranda,M., Nguyen,M., Onodera,C.S., Palm,C.J., Pham,P.K., Quach,H.L., Southwick,A., Tang,C.C., Toriumi,M., Yamada,K., Yamamura, Y., Yu,G., Yu,S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RIKEN Genomic Sciences Center (GSC) members carried out the collection and clustering of RAFL cDNAs (RAFL cDNA : 'RIKEN Arabidopsis Full-Length cDNA'): Seki,M., Narussaka,M., Ishida,J. Satou,M., Kamiya,A., Sakurai,T., Carninci,P., Kawai,J., Hayashizaki,Y. and Shinozaki,K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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ARASGGRFAKKSEVEAGEDAGGRDRERGSATNSSGSEQVETDSNETLNSSGAP"
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Strasbourg 67084, FRANCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NtSET1, a member of a newly identified subgroup of plant SET-domain-containing proteins, is chromatin-associated and its ectopic overexpression inhibits tobacco plant growth plant J. 28 (4), 371-383 (2001)
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/TEARS LA LLON-"MEGGLESGGNIPPSIDKTRYLDVKPLRCLAPVFPSPNGMSSVST
/POSSPFVCVPPTGPFPPGVAPFYPFVAPNDSGREGESQQTPSGYVPNNVFBDSGTGK
KRGRPKVPRARQQABELTPVEVDVEPLLTQLLTSFK LVDLDQVKKADGDKELAGRVLL
VFDLFRRAMTQIDESRDGPGSGRRPLLALASNMLMTKGVEFNQTKRIGNAPGIEVGDIF
FFRHELCLVGLHAPTMAGIDYMSVKLTMDEEPLAVSIVSSGGYDDDGGDGDVLIYTGQ
GGVQRKDGQVFDQKLERGNLALEKSVHRANEVRYIRGVKDVAYPTGKIYITDGLYKIQ
ESMAEKNKYGCNYFKYKLLKYVGGQPGAFKVMKSIQOMKDGVASRVGVILIPDLTSGAES
QPVCLVNDVDDEKGPAYFTYIPSLKYSKFFVMFRPSPSGFHCVGGCQPGDSNCACIJSN
GGFLPYSSLGVLLSYKTLIHECGSACSCPPNCRNRMSQGGFKARLEVFKTKNRGWGLR
SWDPIRGGGFICEYAGEVIDAGNYSDNYIFDATRIYAPLEAERDYNDESRKVPFPDV
SMDPIRGGGFICEYAGEVIDAGNYSDNYIFDATRIYAPLEAERDYNDESRKVPFPDV
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157. .2271
                                                                                                                                                                                                                                                           /protein_id="CAC67503,1"
/db_xref="GI:15485584"
/db_xref="SPTREMBL:Q93YF5"
ISAKNGGNISRFMNHSCSPNVYWQLVVRQSNNEATYHIAFFAIRHIPPMQELTFDYGM
                                                                                                                                                                                                                                                                                                   /product="SET-domain-containing protein"
/protein_id="CAC67503,1"
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                                                                                                                                                                                                                                                                                                                                                                  function="chromatin modeling"
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/note="ecotype: Columbia"
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/db_xref="taxon:3702"
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AB019231 43570 Arabidopsis thaliana genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (24-NOV-1997) Entomology, Oklahoma State University, 127 Noble Research Center, Stillwater, OK 74078, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Palmer, M.J., Guo, 2
Direct Submission
                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ixodid tick, Amblyomma americanum (L.)
Mol. Cell. Endocrinol. 139 (1-2), 45-60 (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Guo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Amblyomma americanum retinoid AF035577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolation of two functional retinoid X receptor subtypes from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Palmer,M.J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Acari;
Parasitiformes; Ixodida; Ixodidae; Amblyomma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amblyomma
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                                                                                                                                                                                                                                                            /codon_start=1
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477 c 659 g 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism-"Amblyomma
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                                                                                                                                                              Score 22;
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0.32;
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DEFINITION

chromosome

5, P1 clone:MGO3

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REFERENCE
AUTHORS
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MEDLINE
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                                                                                                                                                                      CDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein similarity are described as 'unknown protein'.

The software programs used to predict genes include: Grail
(Informatics Group, Oak Ridge National Laboratory,
http://compbio.ornl.gov/Grail-1.3/),
GENSCAN (Chris Burge, MIT, http://CCR-081.mit.edu/GENSCAN.html),
NetGene2 (S.M. Hebsgaard, et al., CES. Technical University of
Denmark, http://www.cbs.dtu.dk/services/NetGene2/) and
SplicePredictor (Volker Brendel, Stanford University,
http://gremlinl.zool.iastate.edu/cgi-bin/sp.cgi).
Genes encoding tRNAs are predicted by tRNAscan-SE
(Sean Eddy, Washington University School of Medicine, St. Louis,
http://genome.wustl.edu/eddy/tRNAscan-SE/).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and Tabata,S.
Structural analysis of Arabidopsis thaliana chromosome 5.
Structural analysis of the regions of 3,076,755 bp covered
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AB019231 BA000015
AB019231.1 GI:3869070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             shorter
The 5' c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        http://www.kazusa.or.jp/kaos/cgi-bin/agd_graph.cgi?c=MGO3
Genes with similarity to proteins in the databases are described in
'product' or 'note' qualifiers. Genes that have no significant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel:81-438-52-3935, Fax:81-438-52-3934)
Address for correspondence: kaos@kazusa.or.jp
For the latest information on annotation of this clone,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (29-OCT-1998) Yasukazu Nakamura, Kazusa DNA Research Institute, Department of Plant Gene Research; 1532-3, Yana, Kisarazu, Chiba 292-0812, Japan (E-mail:ynakamu@kazusa.or.jp,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence features of the regions pl and TAC clones plank Res. 7 (1), 31-63 (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nakamura,Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence may not be the entire insert of this clone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20181125
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                                                                                                                                                                                                                                                                                  join(2732. 2799,3210. 3339,3544. 3693,3907. 4081,

join(2732. 2799,3210. 3339,3544. 3693,3907. 4081,

4399. 4592,4765. 4874,4963. 5062,5271. 5483,5572. 5720,

6096. 6237,6319. 64459,6908. 7096,7473. 7680,7779. 7872,

8102. 8336,8660. 8851,9115. 9213,9389. 9592,9674. 9866,

10082. 10273,10373. 10524,10736. 10891,10971. 11081,

11213. 11307,11608. 11812,11924. 12126,12360. 12411)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              complement(1224. .2102)
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                                                                                                                                                                                                                            KNKDEDVVDEKEKEKLEDEQRSGERKKEKKKKKSDEEIVSEERKSKKKRKSDEEMGS
                                                                                                                                                                                                                                                       DEDVVDEKVKEKLEDEQKSADRKERKKKKSKKNNDEDVVDEKEKLEDEQKSAEIKEKK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unknown
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/clone_lib="Mitsui P1"
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/strain="Columbia"
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                                                                                                                                                                                              EERKSKKKRKLKEIDD"
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /codon_start=1
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/note="gene_id:MGO3.2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein"
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CDS

CDS

VKGQSIDVVVELMKGHRSSLNMADSKGNTALHVATRKGRIKIVELLLDNNETSPSTKA

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EEHPLGEGILORLKEKGGERFRANLSGKEVEFTGERUT ISDDRILKTIEVGITILANGILT
EPECVSRHNIEKLROCVRNGPMKYPGARNVRYPDGSSETLYGOYRKELADELAIGCIV
EPECVSRHNIEKLROCVRNGPMKYPGARNVRYPDGSSETLYGOYRKELADELAIGCIV
EPECVSRHNIEKLROCVRNGPMKYPGARNVRYPDGSSETLYGOYRKELADELAIGCIV
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VPQTEERRATEALTLAMOVONLCTPRKNGELIVASTQDELISSELLTRKDYFYDRAAFSL
ICSYMGDGMDSIDLPTTILKPIELWTGKQIESVLLRRNASIRVYVTLNVKEKKRK
EHGFDETMCINDGWYYFRNSELLSGQLKARILGRGNKDGLYSILIRDYNSHAAAVCMN
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CVGQQTYNGHRARDGFIDRSLPHEPRMSKSPAAKGFVANSFYSGLTATEFFFHTMGGR
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LREFVSLLGVKSASPPOVLYKASGGKYEILFLIYINMQVFKICVFRYEKKIEAGTA
IGTIGAQSIGEBGTOMTLXTHETPAGVASMNITOGVPRIMEIINAKKIESTPVISAELE
NPLELTSARWVKGRIEKTTLGQVAESIEVLMTSTSASVRIILDNKIIEEACLSITPMS
VKNSILKTPRIKLNNDDIRVLDTGGLDITPVVDKSAAFHUHNLKNVLPNIIVNGIKTV
VKNSILKTPRIKLNNDDIRVLDTGGLDITPVTAGDFTMTGCTNANTFOUGKTTOT
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PLTYVLDPNLVLGLFKRMSDKDCELLXIAYRPENLIITCMLVPPLSIRPSVMIGGIQS
NENDLTARLKQIILGNASLHKILSQPTSSPKNMQVWDTVQIEVAAYINSEVRGCQNQP
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vlmysddlkkkligenvikilallkvsaaisfdegvmsclehleavpmsedeeetvvt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                complement(join(12822. .13766,13852. .14406))
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ERVVVAEDMDKSKQIDGKTKWKLFVEGTNLLAVMGTPGINGRTTTSNNVVEVSKTLGI
EAARTTIIDEIGTVMGNHGMSIDIRHMMLLADVMTYRGEVLGIORTGIIKMDKSVLMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             YFNFILDILKCICKRCSNMLLDEKLYEDHLRKMRNPRMEPLKKTELAKAVVKKCSTMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ILMKWFDRFLSKGDDCPNVQRAFEVWWRRAFIRQVLTEPDEFRLQITLY"
complement(17742. .18866)
/note="gb|AAF18598.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IFKLVREEADYDVSRDTLYGLCHRCLTSLVLCLSEVTTOMNDPGKDRGALMGEIAREA
DNMLMMVDILIEKKLCSEFVKLWADQKELANLHSKIPTMYRHEISKITAQICVGIGKG
RILVNRETRFAVLNTWLEALYDDFGWNRRLSSRSLDRKLVEDGLSQTILTLSLRQQQV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /evidence=not_experimental
/protein_id="BAA96934.1"
/db_xref="GI:8777344"
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LLEAAGSSLAAIAKSNGKTALHSAARNGHAEVVKAIVAVEDDTATRTDKKGGTPLHMA
LLEAAGSSLAAIAKSNGKTALHSAARNGHAEVVKAIVAVEDDTATRTDKKGGTPLHMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                 CKSDISKSVVEFRVYNLKREEKRWEEVRDLGDVALFIGDDCSFSVQNPAGDLAGGFIF
YSDYRNGGRSRGICSDGDGVFNVDMQGDFVFSIKPNYFGP"
complement(join(19582 .20451,20673. .21085,21349. .21712)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /translation="mdsssllpsqwsdlpldilelisdrldhdssdtihllclrsvca
Twrlslplsnknnrlskfpkylpfwsssssssgfftlkqsnvykleaplnprtclvkl
Qettpgimryldlfsndricflpenfpskidlqefhyklvrryykmeyanngggavpc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /evidence=not_experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLEELHLPDDSVTLILQRVSSQPSTSSTRTRTDDIFSKLLTGVLQAKDDKARREMKVL
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                                                                                                                                                                                                                                                                                                                                                                                                             /note="gene_id:MGO3.5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNRCIVVDDKGKTVIYDVDFKVSDLAEGLVGGGGHKKHLVECSGGEVFLVDKYVKTVW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FWSLHSDKVVILSSGEDSAIIAIHSGGKLGFLKSGNDEKWKILDNSWNVIYEDIMLYK
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ACCESSION
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AC114102
                                                      SOURCE
                                                                                                                                                                                     DEFINITION
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Best Local Similarity
                              ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                  426 GAATAAAACCATAAAATATTTA 447
                                                                            AC114102
AC114102.2 GI:21732042
HTG; HTGS_PHASE1.
                                                                                                                                                                                  Rattus norvegicus clone CH230-70H23,
                           Norway rat.
Rattus norvegicus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                   AC114102
                                                                                                                                                         ***, 56 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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SDEHRNESGSSGGLSNQDGGSDNGSGTQSSWTKRASDTKSTSPSNQFPDAPNKKGTYE
NGCAHVNRLKEAEDQKEQIGTGSQTGWSMSKKAEEPGDLEKNAKYSVQALERNNDDTL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /translation="MCFNNIETGDEVETERQVFGSSEEDEFRVEDTARNTNNVQISQQ
QQQPLAHVVKWERYLPVRSLKVLLVENDDSTRHIVTALLKNCSYEVTAVPDVLEAWRI
LEDEKSCIDLVLTEVDMPVHSGTGLLSKIMSHKTLKNIPVIMMSSHDSMVLVFKCLSN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          complement(]oin(27357...27393,27488...27555,27625...
27890...27920,28001...28280,28403...28550,28642...(28897...29303,29127...29288,29377...29644))
/note="emb[CAB86035.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HLRLPDDREVADPTAALLPVSPPLPDNCGESFLWRELVNGTINFRDEFYLGKGNFGEV
HRCYESRLNVDGAVKVQNPQNPTGHVEFLAEVTTLRATNHPNVIRLLGNCYGQRNRAI
HYCYESRLNVDGAVKVQNPQNPTGHVEFLAEVTTLRATNHPNVIRLLGNCYGQRNRAI
HYEFAANGDLERHIFAHKTMVGGPRPKGLTLPIRVLDMYTRMKLAVGVAKGLVYLHEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   complement(join(25224. .25460,25563. .25996,26073. .26262,
26345. .26514,26702. .26867))
/note="contains similarity to protein kinase
gene_id:MGO3.7"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VLLTGRKAYDLARPVAKEKLTDWLMPVWTRLEYAPMIVDVALGNKYSIEGLNRLLQTARMCINPQALERPAMDEVETMVREAAAFPVPKEPLVKERISTSK"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EGDDEADDEDPSSIVKFRWNDVVEGTDNFAITHLIGQGTYĞKYYRCKFPEGHKVGÄÄK
IHSNGITGLSETVAETTILKAADHPNVIKLVGKYXGKEKSULVYEFKPEGSLDHHLF
AHARQVQGLTLPTRULDWNTRLRIAVGVAEGILVYVHQGLYAIHRDIKVENILLLDKDEV
PKLTDFGFATKIVYNSDGVERLREFNTRGTQGYIAPEADEFALVSTKSDIYSYGVLLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GEISWLYSTKTDYYSYGAFLLVLFTGRKPFYSDNPAGKKNLTDWFLRVWDRIEDAPYK
GADVALGNRYSVEGLKKIFDTARMCIKAERLERPTMSEVEAMVLEAAAFPVQVPRQVTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product="serine/threonine protein kinase-like"
/protein_id="BAA96937.1"
/db_xref="GI:8777347"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /evidence=not_experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23411. .23541,23892.
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/protein_id="BAA96939.1"
/db_xref="GI:8777349"
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Pred. No. 0.44;
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3, *** SEQUENCING
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TITLE
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Direct Submission
Direct Submission
Submitted (07-MAR-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
3 (bases_1 to 86093)
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Buhay, C., Burcell, K.L., Byrd, N.C., Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Denn, A.L., Dederich, D.A., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwalte, K. J., Draper, H., Dugan, Rocha, S., Durbh, K.J., Davis, D., Dugan, Rocha, S., Durbh, K.J., Davis, D., Dugan, Rocha, S., Durbh, K.J., Davis, D., Davis, Dugan, Rocha, S., Durbh, K.J., Davis, D., Davis, D., Dugan, Rocha, S., Durbh, K.J., Davis, D., Davis, D., Dugan, Rocha, S., Durbh, K.J., Davis, D., Davis, D., Durbh, K.J., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis,
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Submitted (13-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 11, 2002 this sequence version replaced qi:19224429
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Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap: version 0.990329
Consensus quality: 10778 bases at least Q40
Consensus quality: 11488 bases at least Q20
Consensus quality: 12089 bases at least Q20
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arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
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NOTE: This is a 'working draft' sequence. It currently consists of 56 contigs. The true order of the pieces is not known and their order in this sequence record is
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Laboratory, John Innes Centre, Col E-mail: michael.bevan@bbsrc.ac.uk
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                                                                                                         /gene="T24H18_20"
8885. .11005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /translation="MQSKPGRENEEEEVNNHAVQQPMMYAEPWMKNNSFGVVPQARPS GIPSNSSSLDCPNGSESNDVHSASEDGALNGENDGTWKDSQAATSSRSVDNHGMEGND PALSI RNMHDQFLVQPPELVGHY I ACVENPYQDDYYGGLMGAYGHQQLGFRPYLGMPR ERTALPLDMAQEPVYVNAKQYEGILRRRKARAKAELERKVIRDRKPYLHESRHKHAMR RARASGGRFAKKSEVEAGEDAGGRDRERGSATNSSGSEQVETDSNETLNSSGAP" complement (2964...3167)
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4048. .4148.4502. .4778))
/gene="T24H18_10"
/codon_start=1
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4048. .4148,4502. .4778))
/gene="T24H18_10"
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/gene="T24H18_10"
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/gene="T24H18_10"
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/gene="T24H18_10"
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/variety-"Columbia"
/db_xref="taxon:3702"
/chromosome="5"
                                                                                                                                                                                                                                                                                                                                             complement (4048.
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                                         /gene-"T24H18_20"
/note-"Similarity to zinc finger transcription
(PEII), Arabidopsis thaliana, EMBL:AF050463"
                                                                                                                                                                                                                                                                                 complement(4149
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/number="
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/db_xref="GI:7630040"
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/product="zinc finger transcription factor-like protein"
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                          /codon_start=]
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/TIANSLATION-"MASLALSGSCSLAFPLKSRSLSLPRPPSSSLNLTKPLRSLDSRF
/TIANSLATION-"MASLALSGSCSLAFPLKSRSLSLPRPPSSSLNLTKPLRSLDSRF
SLLKSPLPVSLRRRSSTLVKASSTVASASSSPTPPLVPAPVPMGGAAIKPLLASIATG
LILMEYPVPDEGVTRNAWQLLAIFLATIVGIITQPLPLGAVALMGLGASVLTKTLTFAA
AFSAFGDPIPMLTALAFFPARGFIKTGLGNRVAYQFVRLFGSSSLGLGYSLVFSEALL
APAIPSVSARAGGIFLPLVKSLCVACGSNVGDGTEHRLGSWLMLTCFQTSVISSSMFL
TAMAANPLSANLAFNTIKQTIGMTDWAKAAIVPGLVSLIVVPFLLYLIYPPTVKSSPD
APKLAQEKLDKMGPMSKNELIMAATLFLTVGLIMFGAKLGVDAVTAAILGLSVLVTG
VTWKECLAESVAMDTLTWFAALIAMAGYLNKYGLIEMFSQTVVKTVGGLGLSWQLSF
GILVLLYFYTHYFFASGAAHIGAMTTAFLSVSTALGTPYFFASLVLAFGLSLSWGLSTH
GILVLLYFYTHYFFASGAAHIGAMTTAFLSVSTALGTPYFASALVLAFGLSH
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/gene="T24H18_30"
/note="strong similarity to 2-oxoglutarate/malate
translocator precursor, spinach, SWISSPROT:SOT1_SPIOL"
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PGSPSAAQHSTTPPISPSGNGSMPHSSMGMAPAQNNIPALNIPGSNIQLSRLRSSLNARD
IPSEGLSMLHEFEMORGLAGDMHSPREMNHSARRTLNESHEFSAEVASPRRSGD
LAVSSYLSPSHKSALLNQLQNNKQSMLSPIKTNLMSPKNVEQHSLLQQASSPRGGEP
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VSSPTKPHGTDVTFASEKKEYPIDPSLPDIKSGIYSTDEFRMFSFKIRPCSRAYSHDW
TECPFAHPGENARRDPRKFHYTCVPCPDFKKGSCKQGDMCEYAHGVFECWLHPAQYR
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FAADNDVEGFERQLSDVSCINQMGLWYERQREVERMVLEQRTPLMVASLYGSLDVVKF
ILSFPEAELNLSCGPDKSTALHCAASGASVNSLDVVKLLLSVGADPNIPDAHGNRPVD
                                   /translation="MRKPEVAIAASTHQVKKMKKGLWSPEEDSKLMQYMLSNGQGCWS
DVAKNAGLQRCGKSCRLRWINYLRPDLKRGAFSPQEEDLIIRFHSILGNRWSQIAARL
PGRTDNEIKNFWNSTIKKRLKKMSDTSNLINNSSSSPNTASDSSSNSASSLDIKDIIG
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/gene="T24H18_30"
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SDELGRLRKSHSLANNPNREADVSWAQQMLKDSSSPRNGNRVVNMNGARPLTQGGSSV
                                                                                                                                                                                                                                                                        /note="Contains ATP/GTP-binding site motif A (P-loop)
AA51-58; Myb DNA-binding domain repeat signatures
AA23-31; Myb DNA-binding domain repeat signatures AA95
                                                                                                                                                                                                                                                                                                                                                                        complement(join(14756. .15317,16476. .16756))
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                                                                                                                                                                                                                                                                                                                                                                                                                                            complement(join(14756. .15317,16476. .16756))
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /gene="T24H18_30"
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/gene="T24H18_30"
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SFMSLQEQGFVNPSLTHIQTNNPFPTGNMISHPCNDDFTPYVDGIYGVNAGVQGELYF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               complement(12419.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DEFINITION
                                                                                                                                                                                                                                               AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   374 GCAGGAGGAGAGACAGAGAAA 395
Muzny, D.M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbaria, J., Bencon, J., Binage, K., Blankenburg, K., Bonnin, D., Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Bunkett, C., Burrell, K.L., Byrd, N.C., Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J.,
                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                            AC117309
AC117309.3 GI:21745791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rattus
                                                                                                                                                                                                                                                                                          Rattus
                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                           Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                 Norway rat.
                                                                                                                                                                                                                                                                                                                                                                                                         HTG; HTGS_PHASE1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       50 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /translation="MRRSWPTRLILLMVLTVVITMIAAAYGYSSVSSSKHKFPHYKYKAPSPPTTYSPYKYFSPPPVTDSDSAAYVR"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note-"similarity to hydroxyproline-rich glycoprotein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA152-162; UDP-glycosyltransferases signature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94029 bp
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3, *** SEQUENCING
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IN PROGRESS
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Hernandez, O., Hodgson, A., Hogues, M., Holloway, C., Hollins, B., Homsi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Karlsson, E., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C., Karlsson, E., Kelly, S., Khan, U., King, L., Kovar, B., Li, J., Lichtarge, O., Lieu, C., Liu, J., Liu, W., Loulseged, H., Lozado, R. J., Lu, X., Lucler, A., Lucler, R., Luna, R., Ma, J., Lozado, R. J., Lu, X., Lucler, A., Lucler, R., Luna, R., Ma, J., Maheshwari, M., Mapua, P., Martin, R., Martindale, A., Martinez, E., Maheshwari, M., Micker, E., McLeod, M.P., Meador, M., Mei, G., Metzker, M., Miner, G., Mitchell, T., Mohabbat, K., Morgan, M., Morris, S., Moser, M., Nickerson, J., Newtson, N., Nguyen, A., Nguyen, N., Nguyen, N., Nickerson, E., Nwokenkwo, S., Oguh, M., Okwuonu, G., Oragunye, N., Oriedo, R., Partmus, E., Pu, L.L., Quiles, M., Ren, Y., Peters, L., Pickens, R., Primus, E., Pu, L.L., Quiles, M., Ren, Y., Rives, M., Rojas, A., Rojubokan, I., Rolfe, M., Rulz, S., Savery, G., Scherer, S., Scott, G., Shen, H., Shooshtari, N., Sisson, I., Scherer, S., Scott, G., Shen, H., Shooshtari, N., Sisson, I., Sutton, A., Svatek, A., Tabor, P., Tamerisa, A., Tamerisa, N., Thomas, S., Ushani, K., Vasquez, L., Vera, V., Villalon, D., Vinson, R., Wang, Q., Wang, S., Ward-Moore, S., Warren, R., Washington, C., Watlinams, G., Williamson, A., Wleczyk, R., Wooden, S., Worley, K., Weisston, D., Williams, G., Williamson, A., Wheczyk, R., Wooden, S., Worley, K., Weisston, D., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., Williams, G., William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (18-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 14, 2002 this sequence version replaced airanteresce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (10-APR-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gabisi,A., Gao,J., Garcia,A., Garner,T., Garza,N., Gill,R., Gorrell,J.H., Guevara,W., Gunaratne,P., Hale,S., Hamilton,K., Harris,C., Harris,K., Hart,M., Havlak,P., Hawes,A., Hernandez
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             * NOTE: Estimated insert size may differ from sequence length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
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Falls,T., Ferraguto,D., Flagg,N., Ford,J., Foster,P., Frantz,P.,
                                                                                                                                                                                                                                                                 arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                          (see http://www.hgsc.bcm.tmc_edu/docs/Genbank_draft_data.html).
NOTE: This is a 'working draft' sequence. It currently
consists of 50 contigs. The true order of the pieces
is not known and their order in this sequence record is
                                                                                                                                                        as soon as it is available and the accession number will be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap: version 0.990329 Consensus quality: 4555 bases at least Q40 Consensus quality: 49416 bases at least Q30 Consensus quality: 51546 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Center project name: GTRW center clone name: CH230-49C23 ----- Summary Statistics Sequencing vector: Plasmid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
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               \frac{1036}{1136}
1035: contig of 1035 bp in length 1135: gap of unknown length 2169: contig of 1034 bp in length
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REFERENCE AUTHORS

TITLE JOURNAL

COMMENT

JOURNAL REFERENCE AUTHORS TITLE

TITLE

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Unpublished
                                                                                  Direct Submission
Submitted (28 JUN-2001) IBI/MIPS, National Research Center for Environment and Health, Ingolstaedter Landstr. 1, 85758 Neuherberg, Ermail: schoof@gsf.de,kmayer@gsf.de Project Coordinator: Mike Bevan, Molecular Genetics Department, Cambridge Laboratory, John Innes Centre, Colney Lane, NR4 7UJ Norwich, UK, E-mail:
                                          Information on performance of analysis and a more detailed annotation of this entry and other sequences of chromosomes 3, 4 and 5 can be viewed at: http://www.mips.biochem.mpg.de/proj/thal/
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Bevan, M., Murphy, G., Ridley, P., Hudson, S.,
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exon	intron	exon		intron	exon	intron		exon					CDS	gene	gene	exon	intron	exCn							CDS	gene	g ene	
/number=3 complement(1537315451) /gene="At5g12280"	complement(1526515372) /gene="At5g12280"	/Jene="Att5g12280" /number=3	/gene="At5g12280" /number=2 /number=2	/yene- nuyiiio /number-2 complement(1507015141)	complement(1439815069)	complement(143321439/) /gene="At5g12280"	/gene="At5g12280" /number=1	<pre>OFLAQHPGSSTIMVSVPDPDDWKVVKITYQSLSENVASLKEKISGEIQFFTNKQKLRG KAGFIKDNTSLAHYNVGAGEILTLSWW" complement(1409514331)</pre>	DYVLPEGYTIEDYDTIFLTAQIYGRYGEEFWLDLIKEVDNKPQFEFLKPADSKETYFN RLSVVASEGLKRSEKLTSSRMAIVSKAFEYHLRQNEKEBLTHKEGAAKACTLTGALY AFYGPDGCFADIKDEDLFULQHFSAIKNILTKLQPRMPPLGGESEEPEFDESVLVPED	/translation="MVGTLSPDTPDTQFISLCYKLQIVEVVRSPLFSTSRPGGSRGRA /translation="MVGTLSPDTPDTQFISLCYKLQIVEVVRSPLFSTSRPGGSRGRA MRSRLRLHKPIILGQMKFGESDFDVSYDLIFMLEKDNSNEAPAEPPREMRIHMDKIAR VVARKGLRIERKIMNVSEYLAKNQDEGAQEPEAPATLKGDAQHYRRPFLLSDPAPKFP	/product="putative protein" /protein_id="CAC42889.1" /db xref="G1:14586358"		complement(join(1409514331,1439815069,1514215264, 1537315451,1556915717)) 7ene="a+5q17280"	14331,14398 .15717)}	15717 t5g12280"	complement(1288613382) /gene="At5g12270" /number=2	Complement(12/98, 12883) /gene="At5g12270" /number=1		LQILSNDRYRSVEHCVVVNKYCSRVS.HVYEGGFYHDSVLEFLFEVLDANNEMARIKAL VYSDYLKFFFGRPHDGKKTIESIKLP" COMPLEMENT/19212 17971	WPSVCKEQVIDHFKRIKPITERILNILINNLNTIIDESNKETLMGTMRMNFNYYPKCP EPSLAIGTGRHSDINTLTLLLQEDGVLSSLYARATEDGDKWIHVPPIPGAIVVNIGDV	RGEFELPAEFERRYWRGSSVSETAWLTTSENCCIESVLEWRDELKFEYLPQRHDFAAT	/protein_id="CA642888".1" /db_xref="g1:14586357" /t-ss-61=t-on="MotTMTDTTEVPDEVVNOKNGVKGIVDETTITTLDSPYIOPPOER	/product="1-AMINOCYCLOPROPANE-1-CARBOXYLATE OXIDASE-like protein"		to riposing protein	12797 12886		/db_xref="taxon:3702" /chromosome="5" 12212
THICTOR	÷	exon	intron	exon		intron	exon	intron	exon	intron	exon	intron	exon									CDS		gene	gene		exon	intron
90"	/gene="At5912390" /number=6 comol =ment(1758617808)	/number=5 complement(1751317585)	/minimer complement(1743417512) /gene="At5g12290"	complement(1740517433) /gene="a.5912290"	=	/number=4 /number=4 complement(1731917404)	/number=' complement(1719517318) /gene="At5g12290"	/ number = 7 complement(1705417194) /gene="At5g12290"	/	/number=2 complement(1682616942) /gene="At5912290"	/uumber-1 complement(1672516825) /gene="At5g12290"	/number=1 complement(1657416724) /gene="At5g12290"	NSDITITIONAL (1648616573) complement(1648616573) /gene="At5912290"	VENLEVYNKI EARLYKE INNLISSELIKKS ULKKY ZAKOLIVETEKRIMOYOS NFAILAALPAFFISIYMLTVLETWIKKOSKAOGGGRIARIHRRLLVVEIEKRIMOYOS YIEOGRDKDAETVFGLLIYSLERLYRVVEKPARATDEWDFETRSDRVRKATAANIIOA	HEAVEKENI LI WAKI LOGA VOLGA FERLINGUL DANGUL ZELIKA KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KENDENGA SELIKAN KEND	SNLHDIQKSLDEWKSKABUSNAKKAI FALE ERWETER VIDEREN FOR DE DE DE LE COSSSSHMTERMRYLVELEKSALASFLAGLIVELDEKREDLYKIPEKALPSLLAVINGL COSSSSHMTERMRYLVELGSSVDGSYPMPLVFDRLPEVNEEGSQWTDCELTDAINLVGK FSNLEGSFSHLLAVRECDSVDGSYPMPLVFDRLPEVNEEGSQNTENWYHDAKBATMSFESDH	/translation="mdsqppvnestpstalsnegetyppsysylwnrlastlptskpi FLGKISKLYRQTVSRKRSISFPLPLPSDFPSSSTITSNYSADTARIHGVLEEIMADVL FLGKISKLYRQTVSRKRSISFPLPLPSDFPSSSTITSNYSADTARIHGVLEEIMADVL	/product="putative protein" /protein_id="CR642890.1" /db_xref="G1:1458359"	8° tart=1	/gene="At5g12290" /note="similarity to NCA2 protein, yeast, PIR:S54389 /onthins 'Homeobox' domain signature and profile	3659,1883018946,19058. 3719,1991520147))	complement(join(1648616573,1672516825,1694317053, 1719517318,1740517433.1751317585,1780917898,	20147))	complement(join(1648616573,1672516825,1694317053, 1719517318,1740517433,1751317587,1760917898, 1719517318,1740517626,176361788		/gene="At5g12280" /number=5	/number=4 complement(1556915717)	/number=4 complement(1545215568) /gene="At5g12280"

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RESULT 31
AC126575
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                                                                                     Auzny, D.M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alabrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Alabrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Binage, K., Blankenburg, K., Bonnin, D., Bouck, J., Burch, P., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buhay, C., Burch, P., Burkett, C., Burell, K.L., Byrd, N.C., Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Delgado, O., Dann, A.L., Ding, Y., Dinh, H.H., Davila, R., Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Flagg, N., Ford, J., Foster, P., Frantz, P., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gorrell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K., Harris, C., Hollins, B., Hernandez, O., Hodgson, A., Hogues, M., Holloway, C., Hollins, B., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Karlovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L., Li, J., Li, M., Lichtarge, O., Lieu, C., Liu, J., Liu, W., Loulseged, H., Malbeshwari, M., Mapua, P., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin, R., Martin,
Li,J., Li,Z., Lichteryor, Lice,R., Luna,R., Ma,J.,
Lozado,R.J., Lu,X., Lucier,A., Lucler,R., Luna,R., Martinez,E.,
Maheshwari,M., Mapua,P., Martin,R., Martindale,A., Martinez,E.,
Massey,E., Mawhiney,E., McLeod,M.P., Meador,M., Mei,G., Metzker,M.,
Massey,E., Mawhiney,E., McLeod,M.P., Meador,M., Morgan,M., Morris,S.,
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22; Conserv
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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HTG; HTGS_PHASE1.
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/gene="At5g12290"
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/gene="At5g12290"
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/gene="At5g12290"
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/gene="At5g12290"
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Pred. No
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 8;
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Scherer, S., Scott, G., Shen, H., Shooshtari, N., Sisson, I., Sodergren, E., Sonakke, T., Sparks, A., Stanley, H., Stone, H., Sutton, A., Svatek, A., Tabor, P., Tamerisa, A., Tamerisa, K., Tang, H., Tansey, J., Taylor, C., Taylor, T., Telfrod, B., Thomas, N., Thomas, S., Usmani, K., Vasquez, L., Vera, V., Villalon, D., Vinson, R., Wang, Q., Ward-Moore, S., Warren, R., Washington, C., Watlington, S., Williams, G., Williamson, A., Wleczyk, R., Wooden, S., Worley, K., Wu, C., Wu, Y., Wu, Y. F., Zhou, J., Zorrilla, S., Nelson, D., Weinstock, G. and Gibbs, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (24-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (07-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOSET,M., Neal,D., Newtson,J., Newtson,N., Nguyen,A., Nguyen,N., Nguyen,N., Nickerson,E., Nockentwo,S., Oguh,M., Okwuonu,G., Oragunye,N., Oviedo,R., Pace,A., Payton,B., Peery,J., Perez,L., Pickens,R., Primus,E., Pu,L.L., Quiles,M., Ren,Y., Rojas,A., Rojubokan,I., Rolfe,M., Ruiz,S., Savery,G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               * NOTE: Estimated insert size may differ from sequence length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Worley, K.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Direct Submission
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2 (bases 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               runs of N, but the exact sizes of the gaps are unknown.
This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 50 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (bases 1 to 99541)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 55959 bases at least Q40
Consensus quality: 60532 bases at least Q30
Consensus quality: 63142 bases at least Q30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Center project name: GZHJ
Center clone name: CH230-303E15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
Project Information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Center: Baylor College of Medicine Center code: BCM
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1164: gap of
2166: contig
2266: gap of
3327: contig
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AC098517_0
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DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KEYWORDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
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AC098517_1
AC098517_2
AC098517_3
AC098517_3
JS AC098517
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     Muzny, D. M., Adams, C., Adio-Oduola, B., All-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barboroks, S.L., Bankage, R., Blankenburg, K., Bonnin, D., Barbaria, J., Benton, J., Bimage, R., Blankenburg, K., Bonnin, D., Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Bouck, J., Chares, C., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., David, R., Din, H.H., Dederich, D.A., David, R., David, R., David, R., Din, H.H., David, R., Frantz, P., Erantz, P., Erantz, P., Erantz, P., Erantz, R., Garrer, T., Garza, N., Gill, R., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Harris, C., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Homsi, C., Howard, S., Hume, J., Joudah, S., Hennandez, J., Howard, S., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Homsi, F., Howard, S., Hume, J., Joudah, S., Kartovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., J., Li, J., Li, J., Li, J., Li, J., J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li, J., Li,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rattus norvegicus clone CH230-68A4, 171 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rattus norvegicus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AC098517
AC098517.5 GI:21729566
HTG; HTGS_PHASE1.
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/db_xref="taxon:10116"
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20627 c 20564 g 26499 t
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Nickerson, E.,
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85742: gap of unknown 1
90332: contig of 4590
90432: gap of unknown 1
95786: contig of 5354 b
95886: gap of unknown 1
95886: gap of unknown 1
95886: gap of unknown 1
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100.0%; Pred. No. 0.
tive 0; Mismatches
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310000
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  Nwokenkwo, S., Oguh, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 2;
0.48;
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, *** SEQUENCING IN PROGRESS ***,
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On Jul 11, 2002 this sequence version replaced gi:18138579.

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NOTE: This is a 'working draft' sequence. It currently consists of 171 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
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           Submitted (12-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA on Jul 11, 2002 this sequence version replaced gi:17973362.
                                                                                                                                                                                                                                                                                                                                                                                  Submitted (25-NoV-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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IN PROGRESS
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humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk
on Apr 10, 2002 this sequence version replaced gi:19068355.
During sequence assembly data is compared from overlapping clones.
Where differences are found these are annotated as variations
together with a note of the overlapping clone name. Note that the
variation annotation may not be found in the sequence submission
corresponding to the overlapping clone, as we submit sequences with
only a small overlap as described above.
This sequence was finished as follows unless otherwise noted: all
regions were either double-stranded or sequenced with an alternate
chemistry or covered by high quality data (i.e., phred quality >=
30); an attempt was made to resolve all sequencing problems, such
as compressions and repeats; all regions were covered by at least
one plasmid subclone or more than one M13 subclone; and the
assembly was confirmed by restriction digest. The following
abbreviations are used to associate primary accession numbers given
in the feature table with their source databases: Em:, EMBL; Sw:,
SWISSPROT; Tr:, TREMBL; Wp:, WORMPEP; Information on the WORMPEP
http://www.sanger.ac.uk/Projects/C_elegans/wormpep RP23-243J8 is from the RPCI-23 Mouse PAC Library constructed by the group of Pieter de Jong. For further details see http://www.chori.org/bacpac/home.htm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (21-MAY-2002) Wellcome Trust Sanger Institute, Hinxton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AL596180 111914 bp DNA linear ROD 22-MAY-2002 Mouse DNA sequence from clone RP23-243J8 on chromosome 11, complete
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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REFERENCE AUTHORS

(bases 1 to 111914)

JOURNAL

Cambridgeshire, CB10 Direct Submission

lSA, UK. E-mail enquiries:

SOURCE KEYWORDS VERSION ACCESSION DEFINITION

sequence.

AL596180.7 GI:20135751

ORGANISM

Mus musculus

COMMENT

õ

385 AGACAGAGAAAAGAAAAACACA 406

BASE COUNT

29940

þ

/clone\_lib="RPCI-23" 27406 c 26381 g 2 /clone="RP23-243J8"

g

28187

/db\_xref="taxon:10090" /chromosome="11"

/organism="Mus musculus" location/Qualifiers

Query Match Best Local Similarity

3.5%;

Score 22; DB 10; Length 111914; Pred. No. 0.49;

Matches

Conservative

0,

Mismatches

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FEATURES

VECTOR: pBACe3.6.

database can be found at

source

1. .111914

RESULT 33 AL596180

60522 GTGNAAAAAAAAAAAAAAAAA 60543

604 GTGNAAAAAAAAAAAAAAAAA 625

Query Match Best Local Matches

Similarity

Conservative

0;

Mismatches

0

Score 22; Pred. No.

DB 2; 0.48;

length

65778 65878 67076 67176 69031 69131 70707

69030:

gap of unknown length contig of 1198 bp in length gap of unknown length contig of 1855 bp in length

B

97095 AGACAGAGAAAAGAAAAACACA 97116

RESULT

gap of unknown contig of 1576 gap of unknown

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* NOTE: Estimated insert size may differ from sequence length

* (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* consists of 75 contigs. The true order of the pieces

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence

* as soon as it is available and the accession number will

* be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 34683 bases at least 040
Consensus quality: 37389 bases at least Q30
Consensus quality: 39242 bases at least Q20
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REFERENCE AUTHORS KEYWORDS SOURCE

ORGANISM

ACCESSION VERSION DEFINITION

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CELL (DASES 1 CO 115043)

RNS MALTYD, D. M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C.,

RNS Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T.,

Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T.,

Barbaria, J., Benton, J., Bilmags, K., Blankschburg, K., Bonnin, D.,

Bouck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P.,

Bouck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P.,

Bouky, C., Burch, P., Cavazos, S.R., Chacko, J., Chavez, D.,

Cleveland, C.D., Cox, C., Coyle, M., Brown, E., Brown, M., David, R.,

Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A.,

Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D., David, R.,

Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J.,

Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.,

Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M.,

Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M.,

Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M.,

Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M.,

Earnhart, G., Harris, K., Hart, M., Havlak, P., Hale, S., Hamilton, K.,

Gorrell, J.H., Guevara, W., Gunrartne, P., Hale, S., Hamilton, K.,

Harris, C., Harris, K., Hart, M., Havlak, P., Hale, S., Hamilton, K.,

Harris, C., Harris, K., Hart, M., Havlak, P., Hale, S., Hamilton, K.,

Harris, C., Harris, K., Hart, M., Havlak, P., Hale, S., Hamilton, K.,

Gorrell, J.H., Stanker, M., Holloway, C., Hollins, B.,

Homsi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E.,

Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S.,

Kratovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L.,

Kratovic, J., Lichtarge, O., Lieu, J., Liu, W., Loulseged, H.,

Lozado, R.J., Lux, Lucler, A., Lucler, R., Luna, R., Martinez, E.,

Manes, M., Nather, D., Newtson, N., Nuyen, A., Mayren, N.,

Marten, R., Watth, R., Barten, M., Watth, G., Mattinez, E.,

Harris, C., Harris, K., Hartin, R., Stanley, H., Stone, H.,

Peters, L., Pale,
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (17-SEP-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                               Submitted (07-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 6, 2002 this sequence version replaced gi:17940880.
                                                                                                                                                                                                                                                                                                                                                                                                                              Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Worley, K.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Baylor Plaza, Houston, 3 (bases 1 to 115643)
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Direct Submission
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                                HTGS_PHASE1.
                                                                                                                                                  Center code: BCM
Center project name: GACW
                                                                                                                                                                                              Center: Baylor College of Medicine
                                                                                                                                                                                                                                   ---- Genome Center
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JOURNAL REFERENCE AUTHORS TITLE

TITLE

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COMMENT

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* NOTE: Estimated insert size may differ from sequence length

(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* consists of 76 contigs. The true order of the pleces

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 as soon as it is available and the accession number will be preserved.
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----- Summary Statistics
Sequencing vector: Plasmid;
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Rattus norvegicus
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AC123419.2 GI:21902987
                                              Rattus
               HTG; HTGS_PHASE1.
      Rattus norvegicus.
                                                                                                                                           Similarity
                                      us norvegicus clone CH230-248I13, 43 unordered pieces.
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Direct Submission

Submitted (23-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
On Jul 18, 2002 this sequence version replaced gi:21240349.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (29-MAY-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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Direct Submission
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Center project name: GXDX
Center clone name: GXDX
Center clone name: GXDX
Center clone name: GH230-248113
Center clone name: CH230-248113
Center Summary Statistics
Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap: version 0.990329
Consensus quality: 85769 bases at least Q40
Consensus quality: 89524 bases at least Q30
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                                                                                                                                                                                                                                                                                                                  Web site: http://www.hgsc.bcm.tmc.edu/
Contact: hgsc-help@bcm.tmc.edu
Project Information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Center: Baylor College of Medicine Center code: BCM
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NOTE: Estimated insert size may differ from sequence length
(see http://www.bgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently
consists of 43 contigs. The true order of the pieces
is not known and their order in this sequence record is
arbitrary. Gaps between the contigs are represented as
runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          as soon as it is available and the accession number will be preserved.
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          On Apr 22, 2011 this sequence version replaced gi:12329466.

During sequence assembly data is compared from overlapping clones. Where differences are found these are annotated as variations together with a note of the overlapping clone name. Note that the variation annotation may not be found in the sequence submission corresponding to the overlapping clone, as we submit sequences with only a small overlap as described above.

This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one M13 subclone; and the
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22; Conserv
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                            Submitted (20-APR-2001) Sanger Centre, Hinxton, Cambridgeshire CB10 1SA, UK. E-mail enquiries: humquery@sanger.ac.uk Clone
                                                                                                                                                                                                                                                                                                                                AL445433 138918 r
Human DNA sequence from clone
                                                                                                                                                                 requests: clonerequest@sanger.ac.uk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IMPORTANT: This sequence is not the entire insert of clone RP11-234N17 It may be shorter because we sequence overlapping sections only once, except for a 100 base overlap. The true left end of clone RP5-896L10 is at 138819 in this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             http://www.sanger.ac.uk/HGP/Chr1
RP11-234N17 is from the library RPCI-11.1 constructed by the group
of Pleter de Jong. For further details see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      http://www.sanger.ac.uk/Projects/C_elegans/wormpep This sequence was generated from part of bacterial clone contigs of human chromosome 1, constructed by the Sanger Centre Chromosome 1 Mapping Group. Further information can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence. The true right end of clone RP5-846F13 is at 100 in this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SWISSPROT: Tr:, TREMBL; Wp:, WORMPEP; Information on the WORMPEP database can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    abbreviations are used to associate primary accession numbers given in the feature table with their source databases: Em:, EMBL; Sw:,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 http://www.chori.org/bacpac/home.htm
VECTOR: pBACe3.6
                            /note="match: GSS: Em:B98754"
15733. 16309
/note="L2 repeat: matches 2127. .2710 of consensus" 17042. .17208
                                                                                                                                                                                               /note-"MIR repeat: matches 37.
11749. .12081
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="match: GSS: Em:AQ042814"
complement(3711. .3794)
                                                                                                          /note="MER5B repeat: matches 5. .178 of consensus"
                                                                                                                                                14194. .14366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           complement(3711
                                                                                                                                                        /note="L1ME1 repeat: matches 5764. .6118 of consensus"
                                                                                                                                                                                                                                                             /note="MLT1B_repeat: matches 1, .390 of consensus"
                                                                                                                                                                                                                                                                                                            /note="Alujo repeat: matches 1. .310 of consensus"
                                                                                                                                                                                                                                                                                                                                                          /note="Alusq repeat: matches 1. .292 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="match: GSS: Em:AQ519708"
complement(3700..3925)
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complement(3065. .3515)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Sequence confirmed by AC020693 and AC022823 both
sequenced by WIBR."
                                                                                                                                                                                                                                                                                                                                                                                                           'note≃"AluJo repeat: matches 1. .286 of consensus'
                                                                                                                                                                                                                                                                                                                                                                                                                                                               note="MIR repeat: matches 8. ,166 of consensus"
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/db_xref="taxon:9606"
/chromosome="1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note="MIR repeat: matches 19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AIR repeat: matches 81. .145 of consensus"
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24746. .25
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/note="MIR repeat: matches 48. .142
complement(22989. .23434)
complement(2 cc. rm.and55474*
/note="L1MA9 repeat: matches 6037. .6308 of consensus" 34110. .34365
                                                                                                              consensus
                                                                                                                          /note="THE1B-INTERNAL repeat: matches 1.
                                                /note="THE1B repeat: matches 1, .364 of consensus" 33482, .33632
                                                                                                                                                                        /note="THE1B repeat: matches 1. .364 of consensus"
                                                                                                                                                                                                                                                                  'note="L1 repeat: matches 3750. .4361 of consensus"
                                                                                                                                                                                                                                                                                                            /note≖"LlM3a repeat: matches 913. .1541 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                             /note="MIR repeat: matches 12. .151 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                           27905. .28044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="L1MA8 repeat: matches 6006. .6261 of consensus"
27418. .27800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="AluY repeat: matches 81. .306 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="LlMA8 repeat: matches 6261. .6287 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26396. .26836
/note="MLT1C
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20828. .20885
/note="29 copies 2 mer tt 70% conserved"
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18708. .19182
/note="LIMC5 repeat: matches 7447. .7913 of consensus"
                                                                                                                                                                                                                                                                                                                                                        'note="L1M3a repeat: matches 549.
                                                                                                                                                                                                                                                                                                                                                                                                                                           'note="L1MA8 repeat: matches 5924. .6006 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26949.
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18259. .18675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note-"MSTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="L2 repeat: matches 1176. .1564 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="14 copies 2 mer gt 96% conserved"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="L1PA4 repeat: matches 5011. .5403 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'note="L2 repeat: matches 2532. .2750 of consensus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 note="AluJb repeat:
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                                                                                                                                                                                                      JIMA9 repeat: matches 4704. .6037 of consensus 31558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              b repeat: matches 1. .46 of consensus 25839
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       repeat: matches 1. .424 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                repeat: matches 1. .462 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              repeat: matches 1. .111 of consensus"
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                                                                                                                                                                                                                                                                                                                                                   .946 of consensus"
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              Muzny, D. Marie . Metker, M. Lee., Abramzon, S., Adams, C., Alder, J., Allen, C., Allen, H., Alsbrooks, S., Amin, A., Angulano, D., Allen, C., Allen, H., Alsbrooks, S., Amin, A., Angulano, D., Allen, C., Allen, H., Alsbrooks, S., Amin, A., Angulano, D., Anyalebechi, V., Aoyagi, A., Ayodeji, M., Baca, E., Baden, H., Balawin, D., Bandarannaike, D., Barber, M., Barnstead, M., Benahmed, F., Balawin, D., Bandarannaike, D., Barber, M., Barnstead, M., Benahmed, F., Biswalo, K., Blair, J., Blankenburg, K., Blyth, P., Brown, M., Biswalo, K., Blair, J., Blankenburg, K., Blyth, P., Brown, M., Biswalo, K., Blair, J., Blankenburg, K., Blyth, P., Brown, M., Biswalo, K., Blair, J., Barnstead, M., Calderon, E., Cardenas, V., Chen, Z., Chu, J., Chacker, A., Carter, K., Cavazos, I., Ceassar, H., Center, A., Calderon, E., Chan, K., Chen, Y., Chen, Z., Chu, J., Chacker, J., Davis, C., Davy, Carroll, L., De Anda, C., Dederich, D., Davis, C., Davis, C., Davy, Carroll, L., De Anda, C., Dederich, D., Davis, C., Davis, C., Davy, Carroll, L., De Anda, C., Dederich, D., Delgado, O., Denson, S., Dunn, A., Durbin, K., Duval, B., Eaves, K., Egan, A., Escotto, M., Eugene, C., Evans, C.A., Falls, T., Fan, G., Egan, A., Escotto, M., Eugene, C., Evans, C.A., Falls, T., Fan, G., Farser, C., M., Gabisi, A., Ganta, R., Garcia, A., Garner, T., Garza, M., Gebregeorgis, E., Geer, K., Gill, R., Garcia, A., Garner, T., Garza, M., Harnandez, S., Hadlan, S.L., Hodgson, A., Hogues, M., Hernandez, J., Havlak, P., Hawes, A., Henderson, N., Hernandez, J., Hernandez, J., Hadlan, S.L., Hodgson, A., Hogues, M., Marst, C.L., Lebow, H., Levan, J., Idebird, D., Jackson, A., Koyar, C., Karpathy, S., Kelly, S., Kelly, S., Khan, Z., King, L., Koyar, C., Liu, Y., London, P., Longacre, S., Lopez, J., Liu, Y., London, P., Longacre, S., Lopez, J., Liu, Y., London, P., Longacre, S., Lopez, J., Liu, Y., London, P., Longacre, S., Lopez, J., Louiseaed, H., Lozado, R.J., Lu, X., Ma.J.,
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AC114056 DNA linear HTG 24-AUG-
Rattus norvegicus clone CH230-19217, *** SEQUENCING IN PROGRESS
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Lorensuhewa,L., Loulseged,H., Lozado,R.J., Lu,X., Ma,J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="MIR repeat: matches 3. .260 of consensus" 34736. .38491
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38655. .4018
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42450. .42781
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="L1PA8 repeat: matches 3. .3444 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             42508.
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12139. .42417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="L1PA8 repeat: matches 5942. .6163 of consensus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note-"L2 repeat: matches 2647. .2673 of consensus"
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COMMENT

as soon as it is available and the accession number will be preserved.

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1218: contig of 1218 bp in length
1318: gap of unknown length
2614: contig of 1296 bp in length
2714: gap of unknown length
3714: contig of 1000 bp in length
3814: gap of unknown length
5073: contig of 1259 bp in length
5073: gap of unknown length
5173: gap of unknown length

contig

unknown of 1339

JOURNAL

TITLE AUTHORS

REFERENCE

В

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Popovic, D. Primus, E. Pul. L. Puazo, M. Quiroz, J. Racilin, E. Popovic, D. Primus, E. Pul. L. Puazo, M. Quiroz, J. Racilin, E. Reeves, K. Regier, M.A. Reigh, R. Reilly, B. Reilly, M. Ren, Y. Reeves, K. Regier, M.A. Reigh, R. Rives, C. Rodkey, T. Rojas, A. Rose, M. Rose, R. Ruiz, S. J. Sanders, W. Savery, G. Scherer, S. Scott, G. Shatsman, S. Shen, H. Shetty, J. Shvartsbeyn, A. Soott, G. Shatsman, S. Shen, H. Shetty, J. Shvartsbeyn, A. Sors, J. Steinle, M. Stong, R. Sutton, A. Sors, J. Steinle, M. Sodergren, E. Svatek, A. Tabor, P. Taylor, C. Taylor, T. Thomas, N. Thomas, S. Tringey, A. Trejos, Z. Usmani, K. Valas, R. Vera, V. Villasana, D. Walter, B. Wang, J. Wang, O. Wang, S. Warren, J. Walter, B. Wang, J. Wang, S. Warren, J. Wooden, H. Worley, K. Wright, D. Wright, R. Wu, J. Yakub, S. Yen, J. Yoon, L. Yoon, V. Yu, F. Zhang, J. Zhou, J. Zhou, J. Zhou, J. Zhou, J. Zhou, J. Smith, B. R. Holt, R. A. Smith, B. O. Weinstock, G. and Gibbs, R. A.
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A., Perez, A., Perez, L., Pfannkoch, C., Plopper, F., Poindexter, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, A., Perez, 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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Maheshwari,M., Mahindartne,M., Mahmoud,M., Malloy,K., Mangum,A.,
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Direct Submission
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* NOTE: Estimated insert size may differ from sequence length

* (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* NOTE: This is a 'working draft' sequence. It currently

* consists of 80 contigs. The true order of the pieces

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence
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Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 92840 bases at least Q40
Consensus quality: 100722 bases at least Q30
Consensus quality: 105423 bases at least Q20
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TITLE JOURNAL REFERENCE

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Birren, B., Linton, L., Nusbaum, C. and Lander, E. Homo sapiens chromosome 1, clone RP11-76A15
                           Birren, B., Linton, L., Nusbaum, C., Lander, E., Abraham, H., Allen, N., Anderson, S., Baldwin, J., Barna, N., Bastien, V., Beda, F., Boguslavkiy, L., Boukhgalter, B., Brown, A., Burkett, G., Campopiano, A., Castle, A., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Domino, M., Doyle, M., Ferreira, P., Fitzhugh, W., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Goyette, M., Graham, L., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Goyette, M., Graham, L.,
                                                                                                                                                                                                                                       Direct Submission
Submitted (06-FEB-2000) Whitehead Institute/MIT Center Research, 320 Charles Street, Cambridge, MA 02141, USA (bases 1 to 160008)
Galagan,J., Gardyna,S., Ginde,S., Goyette,M., Graham,L.,
Grand-Pierre,N., Grant,G., Hagos,B., Heaford,A., Horton,L.,
                                                                                                                                                                                                                                                                                                                                                                                  Stojanovic, N., Subramanian, A., Talamas, J., Tesfaye, S., Theodore, J., Tirrell, A., Vassiliev, H., Viel, R., Vo, A., Wu, X., Wyman, D., Ye, W.J.,
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Homo sapiens chromosome 1 clone RP11-76A15 map 1, WORKING DRAFT
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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70852: gap of u
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                smit, A.F.A. & Green, P. (1996-1997)
http://ftp.genome.washington.edu/RM/RepeatMasker.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               All repeats were identified using RepeatMasker: Smit, A.F.A. & Green, P. (1996-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (24-AUG-2002) Whitehead Institute/MIT Center for Research, 320 Charles Street, Cambridge, MA 02141, USA On Apr 1, 2000 this sequence version replaced g1:7249264.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Levine,R., Lieu,C., Liu,G., Locke,K., Macdonald,P., Marquis,N., McCarthy,M., McEwan,P., McGurk,A., McKernan,K., McPheeters,R., Meldrim,J., Meneus,L., Mihova,T., Miranda,C., Mlenga,V., Morrow
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NOTE: This is a 'working draft' sequence. It currently consists of 38 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Quality coverage: 3.2 in Q20 bases; Quality coverage: 3.3 in Q20 bases;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequencing vector: M13; M77815; 100% of reads Chemistry: Dye-terminator Big Dye; 100% of reads Assembly program: Phrap; version 0.960731 Consensus quality: 132595 bases at least Q40 Consensus quality: 145388 bases at least Q30 Consensus quality: 151459 bases at least Q20 Consensus quality: 151459 bases at least Q20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Center: Whitehead Institute/ MIT Center for Genome Research
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Web site: http://www-seq.wi.mit.edu
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Insert size: 156308; sum-of-contigs
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12988 15511: con
15512 15611: gap of
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                              118: gap of 100 bp
20410: contig of 2592 bp in length
110: gap of 100 bp
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79: gap of 100 bp
12887: contig of 1808 bp in length
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ll be updated with the finished sequence
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contig of 1689 bp in length
of 100 bp
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131971 142739: cont
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120993 131870: cont
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31661 35351: contig of 3891 bp in length
35352 35451: gap of 100 bp
35452 38476: contig of 3025 bp in length
38477 38576: gap of 100 bp
38577 41099: contig of 2523 bp in length
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26905: gap of 100 br
29287: cont'
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58982: contig of 4099 bp in length
100 bp
63451: contig of 4369 bp in length
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187: gap of 100 bp
31560: contig of 2173 bp in length
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contig of 3974 bp
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contig of 10769 bp in length
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104 bp in 7
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REFERENCE
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AC020693
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JOURNAL
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Best Local
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RS Birren, B., Linton, L., Nusbaum, C., Lander, E., Abraham, H., Allen, N., Anderson, S., Baldwin, J., Barna, N., Bastien, V., Beda, F., Anderson, S., Baldwin, J., Barna, N., Bastien, V., Beda, F., Boukslavkiy, L., Boukhgalter, B., Brown, A., Burkett, G., Callymore, A., Castle, A., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Domino, M., Doyle, M., Ferreira, P., Fitzhugh, M., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Goyette, M., Graham, L., Gand-Pierre, N., Graham, L., Johnson, R., Jones, C., Kann, L., Karatas, A., Howland, J.C., Iliev, I., Johnson, R., Jones, C., Kann, L., Karatas, A., Kiein, J., Laroque, K., Lanazares, R., Landers, T., Lehoczky, J., Levine, R., Lieu, C., Liu, G., Locke, K., Macdonald, P., Marquis, N., McCarthy, M., McEwan, P., McGurk, A., McKernan, K., McPheeters, R., Meldrim, J., Weneus, L., Mihova, T., Miranda, C., Mlenga, V., Morrow, J., Murphy, T., Naylor, J., Norman, C.H., O'Connor, T., O'Donnell, P., O'Neil, D., Olivar, T. M., Oliver, J., Peterson, K., Pierre, N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
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BoguslavKiy, L., Boukhgalter, B., Brown, A., Burkett, G., Castle, A.,
Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P.,
Pearellano, K., Dewar, K., Domino, M., Doyle, M., Fenestor, J.,
Ferreira, P., FitzHugh, W., Forrest, C., Gage, D., Galagan, J.,
Gardyna, S., Grant, G., Hagos, B., Heaford, A., Horton, L.,
Gardyna, S., Grant, G., Hagos, B., Heaford, A., Horton, L.,
Howland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J.,
Howland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J.,
Howland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J.,
Howland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J.,
Macdonald, P., Marquis, N., McEwan, P., McGurk, A., McKernan, K.,
McDheeters, R., Meldrim, J., Meneus, L., Morrow, J., Naylor, J.,
Morman, C.H., O'Connor, T., O'Donnell, P., Olivar, T.M., Peterson, K.,
Pierre, N., Pisani, C., Pollara, V., Raymond, C., Riley, R., Rothman, D.,
Rova, A., Santoa, B., Severy D., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Santoa, B., Sa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Roy,A., Santos,R., Severy,P., Spencer,B., Stange-Thomann,N., Stojanovic,N., Subramanian,A., Talamas,J., Tesfaye,S., Theodore,J. Tirrell,A., Vassiliev,H., Viel,R., Vo,A., Wu,X., Wyman,D., Ye,W.J. Zimmer,A. and Zody,M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (08-JAN-2000) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 163932)
Birren,B., Linton,L., Nusbaum,C. and Lander,E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Birren, B., Linton, L., Nusbaum, C., Lander, E., Abraham, H., Allen, N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AC020693.3 GI:7249430
HTG; HTGS_PHASE1; HTGS_DRAFT.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE, 8 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AC020693
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (bases 1 to 163932)
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15612. .17718
/note="assembly_fragment"
17819. .20410
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9442. .10979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12988.
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11080. .12887
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 160008;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear HTG 24-AUG-24
9 map 6, WORKING DRAFT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Theodore, J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24 - AUG - 2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMMENT
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misc_feature
                                                                         misc_feature
                                                                                                                                           misc_feature
                                                                                                                                                                                                                                                                                   misc_feature
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* This record will be updated with the finished sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (24-AUG-2002) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On Mar 16, 2000 this sequence version replaced gi:6730833. All repeats were identified using RepeatMasker: Smit, A.F.A. & Green, P. (1996-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pisani,C., Pollara,V., Raymond,C., Riley,R., Rogov,P., Rothman,D., Roy,A., Santos,R., Schauer,S., Severy,P., Spencer,B., Stange-Thomann,N., Stojanovic,N., Subramanian,A., Talamas,J., Tesfaye,S., Theodore,J., Tirrell,A., Travers,M., Trigillo,J., Vassillev,H., Viel,R., Vo,A., Wilson,B., Wu,X., Wyman,D., Ye,W.J., Vassillev,H., Viel,R., Vo,A., Wilson,B., Wu,X., Wyman,D., Ye,W.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          http://ftp.genome.washington.edu/RM/RepeatMasker.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2756: contig of 2756 bp in length
2757 2856: gap of
2857 7124: contig of 4268 bp in length
7125 7224: gap of
100 bp
7225 14958: contig of 734 bp in length
14959 15058: gap of
100 bp
15059 21054: contig of 5996 bp in length
21055 21154: gap of
21055 21154: gap of
21155 31520: contig of 100 bp
31521 31630: gap of
31521 31630: gap of
31521 31630: contig of 10366
31521 31630: gap of
31521 31630: gap of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequencing vector: M13; M77815; 100% of reads Chemistry: Dye-terminator Big Dye; 100% of reads Assembly program: Phrap; version 0.960731 Consensus quality: 159231 bases at least Q40 Consensus quality: 161639 bases at least Q20 Consensus quality: 162468 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     58638 58737: gap of 100 bp
58738 110374: contig of 51637 b
110375 110474: gap of 100 bp
110475 163932: contig of 53458 k
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Insert size: 160000; agarose-fp
Insert size: 16323; sum-of-contigs
Quality coverage: 5.4 in Q20 bases; agarose-fp
Quality coverage: 5.2 in Q20 bases; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Center project name: L5272
Center clone name: 574_H_19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: sequence_submissions@genome.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Web site: http://www-seq.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Center code: WIBR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Center: Whitehead Institute/ MIT Center for Genome Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submission
                                                                                                                                               2857.
                                                                                                                                                                                                             clone_end:T7
                                                                                                                                                                     vector_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1. .163932
                                                                  7225. .14958
                                                                                                                                                                                                                                                                                                             /clone="RP11-574H19"
/clone_lib="RPCI-11 Human Male BAC"
                         /note="assembly_fragment"
                                                                                               /note="assembly_fragment"
                                                                                                                                                                                                                                          /note="assembly_fragment
                                                                                                                                                                                                                                                                                                                                                                                                                /chromosome="6"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zainoun, J., Zimmer, A. and Zody, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1737: gap of 100 bp 110374: contig of 51637 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ·- Genome Center
.21054
                                                                                                                                                                        side:right"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bp in length
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Muzny, D.M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Bimage, K., Blankenburg, K., Bonnin, D., Bouck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Bouk, J., Bowle, S., Brieva, M., Brown, E., Brydn, M.C., Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Denn, A.L., Ding, Y., Dinh, H.H., Delance, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delance, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delance, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delance, K.R., Carroll, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gabisi, A., Gao, J., Hart, M., Havlak, P., Habes, A., Hernandez, J., Hernandez, J., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E., Jacobson, E., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C., Karlsson, E., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C., Karlsson, E., Kelly, S., Khan, U., King, L., Lucier, R., Luna, R., Ma, J., Lozado, R.J., Lu, X., Lucier, A., Lucier, R., Luna, R., Ma, J., Lozado, R.J., Lux, Lucier, A., Lucier, R., Luna, R., Ma, J., Mahshan, M., McLeod, M.P., Meador, M., Mei, G., Metzker, M., Michell, T., Mohabbat, K., Morgan, M., Morris, S., Michell, T., Mohabbat, K., Morgan, M., Morris, S.,
                                                                                Miner,G., Miner,Z., Mitchell,T., Mohabbat,K., Morgan,M., Morris,S., Moser,M., Neal,D., Newtson,J., Newtson,N., Nguyen,A., Nguyen,N., Nguyen,N., Nickerson,E., Nwokenkwo,S., Oguh,M., Okwuonu,G., Oragunye,N., Oviedo,R., Pace,A., Payton,B., Peery,J., Perez,L., Peters,L., Pimus,E., Pu,L.L., Quiles,M., Ren,Y., Rives,M., Rojas,A., Rojubokan,I., Rolfe,M., Ruiz,S., Savery,G., Scherer,S., Scott,G., Shen,H., Shooshtari,N., Sisson,I., Sodergren,E., Sonake,T., Sparks,A., Stanley,H., Stone,H., Shooshtari, N., Stone,H., Sodergren,E., Sonake,T., Sparks,A., Stanley,H., Stone,H., Scott, S., Sonake,T., Sparks,A., Stanley,H., Stone,H., Shooshtari, N., Stanley,H., Stone,H., Sodergren,E., Sonake,T., Sparks,A., Stanley,H., Stone,H., Shooshtari, N., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stone,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,H., Stanley,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
Sutton,A., Svatek,A., Tabor,P., Tamerisa,A., Tamerisa,K., Tang,H., Tansey,J., Taylor,C., Taylor,T., Telfrod,B., Thomas,N., Thomas,S., Usmani,K., Vasquez,L., Vera,V., Villalon,D., Vinson,R., Wang,Q.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  194402 bp DNA linear HTG 11-JUL-2002 us norvegicus clone CH230-10H22, *** SEQUENCING IN PROGRESS 77 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="assembly_fragment"
21155. .31520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="assembly_fragment"
110475. .163932
/note="assembly_fragment"
, 30930 c 30043 g 50325
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31621. .58637
/note="assembly_fragment"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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JOURNAL
REFERENCE
AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE
AUTHORS
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Submitted (17-SEP-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plazz, Houston, TX 77030, USA
3 (bases 1 to 194402)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (11-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Baylor Plaza, Houston, TX 77030, USA on Jul 10, 2002 this sequence version replaced gi:17943838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 77 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               as soon as it is available and the accession number will be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329
Consensus quality: 127428 bases at least Q40
Consensus quality: 136154 bases at least Q30
Consensus quality: 142855 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Center project name: GENS
Center clone name: CH230-10H22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Center: Baylor College of Medicine
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10893
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12126
12226
13687
13787
          15000
16353
16453
17992
18092
19397
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5771
5871
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2824: gap of unknown length
4385: contig of 1561 bp in length
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                                                                                                     gap of contig gap of contig gap of contig gap of contig gap of contig
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contig
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                                                                      gap of
          f unknown
y of 1305
unknown
                                                                                                        unknown
of 1539
                                                                                                                                                                     of 1113
unknown
of 1353
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of 1011
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                                                                                                                                                                                                                                                                      unknown
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                                                                                                                                                                                                                                                                                                                                                                                                      unknown
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length
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ACCESSION VERSION

AC096152.4 GI:21723293

HTGS\_PHASE1.

DEFINITION AC096152/c

Rattus

SOURCE KEYWORDS

ORGANISM

Rattus norvegicus Norway rat

REFERENCE

Rattus

(bases 1 to 194402)

AUTHORS

BASE COUNT ORIGIN

misc\_feature

misc\_feature

vector\_side:left" 58738. .110374

clone\_end:SP6

misc\_feature

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Query Match Best Local Similarity

Matches

Conservative

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Db 124291 GAAAAACACAGCATGAGAACAC 124312

397 GAAAAACACAGCATGAGAACAC 418

Wang,S., Ward-Moore,S., Warren,R., Washington,C., Watlington,S., Williams,G., Williamson,A., Wleczyk,R., Wooden,S., Worley,K., Wu,C., Wu,Y., Wu,Y.F., Zhou,J., Zorrilla,S., Nelson,D., Weinstock,G. and Gibbs,R.

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RESULT 42
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ORGANISM
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                                                                                                        Alabrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Binage, K., Blankenburg, K., Bonks, T., Barbaria, J., Benton, J., Binage, K., Blankenburg, K., Bonks, T., Burkett, C., Burch, H., Bryant, N.P., Burkett, C., Burch, E., Brown, M., Bryant, N.P., Bunks, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Bunks, J., Bowie, S., R., Chacko, J., Chavez, D., Chen, Z., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwalte, K.J., Drager, H., Dugan-Rocha, S., Durbin, K.J., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Earnhart, C., Edgar, D., Flagg, N., Ford, J., Foster, P., Frantz, P., Flagg, N., Ford, J., Foster, P., Frantz, P., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gurell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K., Harris, C., Harris, K., Hart, M., Haylak, P., Hawes, A., Hernandez, J., Homsi, F., Howard, S., Lucier, R., Julyk, S., Hume, J., Jackson, L., Katlsson, E., Kallys, S., Khan, U., King, L., Korvah, J., Kovar, C., Kratovic, J., Kureshi, A., Landry, N., Ledi, B., Lewis, L.C., Lewis, L.C., Liu, Y., Lichtarge, O., Lieu, C., Liu, J., Liu, W., Loulseged, H., Lozado, R.J., Lu, X., Lucier, R., Luna, R., Martinez, E., Massey, E., Mavhiney, E., McLeod, N.P., Meadonri, M., Morris, S., Massey, E., Mayliner, G., Miner, Z., Mitchell, T., Mohabbat, K., Morgan, M., Morris, S., Massey, E., McCheod, N.P., Paeca, A., Payton, B., Peery, J., Perez, L., Pickens, R., Primus, E., Put, L.I., Qulles, M., Ren, Y., Rives, M., Rojas, A., Rojubokan, I., Rolfe, M., Ruiz, S., Savery, G., Scherer, S., Scott, G., Shen, H., Shooshtari, N., Sisson, I., Sodergen, S., Watches, A., Tabor, P., Tamerisa, A., Tamerisa, A., Tang, H., Mang, O., Wall, J., Watchell, S., Kelson, D., Warsh, R., Wang, O., Wall, J., Watchell, S., Kelson, D., Warsh, R., Wang, O., Wall, J., Watchell, S., Kelson, D., Watchell, S., Kelson, D.
Unpublished 2 (bases 1
                                                                            Wu,C., Wu,Y., Wu,Y.F., Zhou,J., Zorrilla,S., Nelson,D., Weinstock,G. and Gibbs,R.
                                                     Direct Submission
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (23-JAN-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, Tx 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 1(Assembly program: Phrap; version 0.9);
Consensus quality: 153108 bases at 16;
Consensus quality: 159647 bases at 16;
Consensus quality: 166406 bases at 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct Submission
Submitted (13-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Worley, K.C.
Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 62 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                     arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                              as soon as it is available and the accession number will be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
Project Information
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                                                                                                                                                                                                                               Birren, B., Nusbaum, C., Lander, E., Ali, A., Allen, N., Anderson, S., Barna, N., Bastien, V., Bloom, T., Boguslavkiy, L., Boukhgalter, B., Camarata, J., Chang, J., Chazaro, B., Choepel, Y., Collymore, A., Cook, A., Cook, P., DeArellano, K., Dewar, K., Dlaz, J.S., Dodge, S., Faro, S., Ferreira, P., FitzGerald, M., Gage, D., Galagan, J., Gardyna, S., Gord, S., Graham, L., Grand-Pierre, N., Hagos, B., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamata, A., Kells, C., Landers, T., Levine, R., Lindblad-Toh, K., Liu, G., MacCean, C., Macdonald, P., Major, J., Matthews, C., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., Murphy, T., Naylor, J., Nenens, C., Norman, C., Rogov, P., Murphy, T., Naylor, J., Nenens, J., Peterson, K., Phunkhang, P., Pierre, N., Raymond, C., Retta, R., Rise, C., Rogov, P., Roman, J., Roy, A., Schauer, S., Schupback, R., Seaman, S., Severy, P., Smith, C., Spencer, B., Stange-Thomann, N., Stojanovic, N., Talamas, J., Viel, R., Vo, A., Wilson, B., Wu, X., Wyman, D., Young, G., Zainoun, J., Zembek, I., Zimmer, A. and Zodv, M.
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Direct Submission
L. Submitted (23-NOV-2001) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA
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                                            Submitted (21-AUG-2002) Whitehead Institute/MIT Center for Genome
Research, 320 Charles Street, Cambridge, MA 02141,
                                                                                                                                                     Direct Submission
                                                                                                                                                                                                                Zembek, L., Zimmer, A. and Zody, M.
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Birren,B., Nusbaum,C. and Lander,E.
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Mus musculus clone RP23-334D14,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stojanovic, N.,
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* NOTE: This is a 'working draft' sequence. It currently consists of 25 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chemistry: Dye-terminator Big Dye; 100% of re Assembly program: Phrap; version 0.960731 Consensus quality: 191519 bases at least Q40 Consensus quality: 194846 bases at least Q30 Consensus quality: 195925 bases at least Q20 Consensus quality: 195925 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Quality coverage: 5.8 in Q20 bases; agarose-fp Quality coverage: 5.8 in Q20 bases; sum-of-contigs
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Center clone name: 334_D_14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Center code: WIBR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6543 8314: contig of 1772 bp in length

8315 8414: gap of 100 bp

8415 1054: contig of 2540 bp in length

10955 11054: gap of 100 bp

11055 14504: contig of 3450 bp in length

14505 14604: gap of 100 bp

14605 1981: contig of 5277 bp in length
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Insert size: 196443; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequencing vector: Plasmid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ----- Summary Statistics
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135934: gap of 
151141: cont
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135834: contig of 36621 bp in length
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                                                                                                                                                                                         256: gap of 100 bp
63272: contig of 10016 bp in length
372: gap of 100 bp
73192: contig of 9820 bp in length
                                                                                                106: gap of 100 bp
99113: contig of 13007 bp in length
                                                                                                                                               100 bp
86006: contig of 12714 bp
                                                                                                                                                                                                                                                                                               374: gap of 100 bp
53156: contig of 8282 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24662:
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44774: contig of 7723 bp in length
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30542: contig of 5780 bp in length
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0: gap of 100 bp
6442: contig of 1682 bp in length
2: gap of 100 bp
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contig of 15207 bp in length
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162189 162288: gap of 100 bp
162289 175380: contig of 14092 bp in length
176381 176480: gap of 100 bp
176481 187921: contig of 11441 bp in length
187922 198021: gap of 100 bp
188022 198043: contig of 10822 bp in length.
Location/Qualifiers
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37052. .44774
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1904. .3084
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/clone="RP23-334D14"
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Bryant, N., Buhay, C., Burch, P., Burrell, K., Calderon, E.,
Cardenas, V., Carter, K., Cavazos, I., Ceasar, H., Center, A.,
Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chen, Z., Chen, Z.,
Cleveland, C., Cockrell, R., Cox, C., Coyle, M., Cree, A., D'Souza, I.,
Davila, M.L., Davis, C., Davy, Carroll, L., De Anda, C., Dederich, D.,
Delgado, O., Denson, S., Deramo, C., Ding, Y., Dinh, H., Divya, K.,
Derper, H., Dugan-Rocha, S., Dunn, A., Durbin, K., Daval, B., Eaves, K.,
Egan, A., Escotto, M., Eugene, C., Evans, C.A., Falls, T., Fan, G.,
Fernandez, S., Finley, M., Flagg, N., Forbes, L., Foster, M., Gourara, M.,
Gebregeorgis, E., Geer, K., Gill, R., Garda, A., Garrer, T., Garza, M.,
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Newton, W., Norris, S., Nwackelemeh, O., Okwuonu, G.,
Olarpunsagoon, A., Pal, S., Parks, K., Pasternak, S., Poul, H.,
Reeves, K., Regler, M., Reigh, R., Reilly, B., Reilly, M., Ren, Y.,
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Song, X., Soretle, R., Sosa, J., Steinle, M., Stern, S.,
Song, A., Trejos, Z., Usmani, K., Valas, R., Vera, V., Villasana, D.,
Walson, H., Wal
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Biswalo,K., Blair,J., Blankenburg,K., Blyth,P., Brown,M.,
Bryant,N., Buhay,C., Burch,P., Burrell,K., Calderon,E.,
Submitted (29-MAY-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 206648)
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*** SEQUENCING IN PROGRESS ***,
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COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              * NOTE: Estimated insert size may differ from sequence length

(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* consists of 94 contigs. The true order of the pieces

is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence

* as soon as it is available and the accession number will

* be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rat Genome Sequencing Consortium. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequencing vector: Plasmid; Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap: version 0.990329 Consensus quality: 135401 bases at least 040 Consensus quality: 143073 bases at least 030 Consensus quality: 143075 bases at least 020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Center project Information Center project name: GLJV Center clone name: CH230-68C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
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SOURCE KEYWORDS VERSION

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                                                                                                                                                                                                                                                                                                     Submitted (10-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 9, 2002 this sequence version replaced gi:17941347.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (15-SEP-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 209095)
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                      Center clone name: CH230-4B10
                                                                                                                                       Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                              Center: Baylor College of Medicine Center code: BCM
                                                                                                                                                                                                                                                                                          ----- Genome Center
Summary Statistics
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* NOTE: Estimated insert size may differ from sequence length

* (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* consists of 50 contigs. The true order of the places

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence

* as soon as it is available and the accession number will

* be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequencing vector: Plasmid; chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329 Consensus quality: 167289 bases at least Q40 Consensus quality: 173510 bases at least Q30 Consensus quality: 178689 bases at least Q30
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Search completed: November Job time : 3364.8 secs
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COMMENT

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## SUMMARIES

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353.4	354.4	354.4	360.6	363.8	365.4	Score
95.8	96.0	96.0	97.7	98.6	99.0	Query Match 1
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FEATURES

High quality sequence start: 4 High quality sequence stop: 702.

Location/Qualifiers

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REFERENCE AUTHORS DEFINITION ORGANISM JOURNAL Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

CE 1 (bases 1 to 704)

RS NIH-MGC http://mgc.nci.nih.gov/.

RS NIHonal Institutes of Health, Mammalian Gene Collection (MGC)

AL Unpublished (1999)

Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov

Tissue Procurement: Life Technologies, Inc.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

http://image.llnl.gov
plate: LLAMI1618 row: a column: 21 mRNA sequence. BI919074 B1919074 704 bp mRNA linear EST 16-OCT-2001 503180881F1 NIH\_MGC\_121 Homo sapiens cDNA clone IMAGE:5244956 5', EST BI919074.1 GI:16200128 human.

SOURCE KEYWORDS VERSION ACCESSION RESULT 1 BI919074

ALIGNMENTS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              366;
                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                 NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                             Contact: Robert Strausberg, Ph.D
                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                    36 DNA Sequence 10. Jul-2002 mRNA linear EST 15-JUL-2002 AGENCOURT_8301981 NIH_MGC_102 Homo sapiens cDNA clone IMAGE:62751215', mRNA sequence
                                                                                                                                                                                                                                                                          BQ668420.1 GI:21778667
                                                                                                                                                                                                                                                                                              BQ668420
cDNA Library Preparation: Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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/clone="IMAGE:5244956"
/clone_lib="NIH_MGC_121"
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 747)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                  603181571F1 NIH_MGC_121 Homo sapiens cDNA clone IMAGE:5245648 5',
                                                                                                                  Homo sapiens
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                                                                                                                                        human.
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/lab_host="DH10B (phage-resistant)"
/note="Organ: sallvary gland; Vector: pOTB7; Site_1: xhoI;
Site_2: ECORI; CDNA made by Oligo-dT priming.
Directionally cloned into EcORI/xhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-CDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."

96 a 314 c 285 g 191 t 1 others
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/db_xref="taxon:9606"
/clone="IMAGE:6275121"
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Pred. No. 3.2e-102;
0; Mismatches 4;
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mRNA sequence.
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BG753617.1 GI:14064270
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11619 row: n column: 17
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Tissue Procurement: Life Technologies, Inc.
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Contact: Robert St
                                                                       BG753617 670 bp r 602732803F1 NIH_MGC_43 Homo sapiens
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/db_xref="taxon:9606"
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/clone_lib="NIH_MGC_121"
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98.4%;
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                                                                                                                                                                                                                  GGCAACTTTTTGCGGATTGTTCTTGCTTNCAGGCTTTGCGCTGCAAATCCAGTGCTACCA
                                                                                                                                                                  CTGCTCCCCAGGG-AAACTGAACTCAGTTTGCATCAGCTGCTGCTGCAACACCCCCTCTTTGTA 299
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                                                                                       ACGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCA 359
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                                                                        ACGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCAGGCCAGGGCTCCGCA
                                                                                                                                             CTGCTCCCCAGGGCAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTA
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Tissue Procurement: ATCC
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1. (bases 1 to 670)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene
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//tissue_type="normal pigmented retinal epithelium"
//lab_bost="DH10B (phage=resistant)"
//note="Organ: eye: Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; CDNA made by Oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5; adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library. | "
401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
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                 CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCAT-GGCTCCGCA
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                                                                                                                                                                                                                                                                                                                        GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
                                                                                            CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
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CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCAGGCCATGGGCTCCGGA
                                                                         CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTTTTGTAA
                                                                                                                                                CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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NIH-MGC http://mgc.nci.nih.gov/.
National institutes of Health, M
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603180565F1 NIH_MGC_121
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cDNA Library Arrayed by: The I.M.A.G.E. CONSORTIUM (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://imagge.llnl.gov
Plate: LLAM11617 row: a column: 18
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/db_xref="taxon:9606"
/clone="ib-MAGE:5244569"
/clone_lib-"NIH_MGC_121"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note-"organ: brain; Vector: pCMV-SPORT6; Site_1: Not1; Site_2: ECORV (destroyed); RNA source anonymous pool of 3 fetal brains, female age 20 weeks, female age 24 weeks, and male age 26 weeks. Library is oligo-dT primed and directionally cloned (ECORV site is destroyed upon cloning). Average insert size 1.7 kb, insert size range 0.7-3.5 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 017. Note: this is a NIH MGC Library."
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                                 CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAA
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BP 191 91006 EVRY cedex - France
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/clone_lib="LTI_FL013_FBrn1"
/clone_lib="LTI_FL013_FBrn1"
/dev_stage="pooled tissue from post conception fetuses (20 week, 24 week and 26 week)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                         pcMvSPORT 6 vector. Library was constructed by Life Technologies. Contact: Feng Liang Life Technologies, a division of Invitrogen 9800 Medical Center Drive Rockville Maryland 20850, USA Fax: (1) 301 610 8371 Email: fliang@lifetech.com URL: http://fulllength.invitrogen.com" 7 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note-"Organ: Fetal brain; Vector: pCMVSPORT 6; 1st strand CDNA was primed with a NotI-Oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and Eco RV sites of the and Colomed into the Not I and Eco RV sites of the
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                    GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
TCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCAGCACCCCCCTCTTTGTA
                                       TCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTA 299
                                                                                                      ACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCATCGCCTCTGCCCGGGTACCAGTCCT
                                                                                                                                                                   GGTGAACGTTCAAGACATGTGTCAGAAACGAAGTGATGGAGCAAAGTGCCGGGATCATGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis, M.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
Plate: LLAM11784 row: d column: 08
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National Institutes of Health, Mammalian
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Site_2: SalI; Cloned unidirectionally, Primer: Oligo
Average insert size 2.3 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
a 231 c 204 g 132 t
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/db_xref="taxon:9606"
/clone="IMAGE:5309527"
/clone_lib="NCI_CGAP_Brn67"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab_host="DH10B (T1 phage-resistant)"
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98.6%;
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                                                                             GGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA
CGGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGT
                                     CGGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGT 179
                                                                                              -GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tissue Procurement: Miklos Palkovits, M.D., Ph.D. cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki Toshiyuki and Piero Carninci (RIKEN) cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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BI669845
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603293440F1 NIH_MGC_96 Homo sapiens
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National Institutes of Health, Mammalian Gene Collection (MGC)
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Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can
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                                                                                                                                                                                                                                                                                                                                    /note-"Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI, Site_2: SalI-XhoI (gtcga); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library." a 231 c 202 g 130 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="IMAGE:5312889"
/clone_lib="NIH_MGC_96"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="hypothalamus"
/lab_host="DH10B"
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287 AGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA 346
                                                                                                                                                           227
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                                    60 AGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA 119
                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                               1 GGCAACTTTTTGCGGATTGTTCTTGCTTNC-AGGCTTTGCGCTGCAAATCCAGTGCTACC 59
                                                                                                                                                           GGCAACTTTTTGCGGATTGTTCTTGCTTCCAAGGCTTTGCGCTGCAAATCCAGTGCTACC 286
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Tlasue Procurement: David N. Louis, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 1010)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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BF345141
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Plate: LLAM9422 row: d column: 05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="DH10B (T1 phage-resistant)"
//note="Organ: brain; Vector: pCMV-SPORT6; Site_1: Not1;
//note="Organ: brain; Vector: pCMV-SPORT6; Site_1: Oligo dT.
Site_2: Sall; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 2.3 kb. Constructed by Life
Average insert size 2.3 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
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/db_xref="taxon:9606"
/clone="IMAGE:4153684"
/clone_lib="NCI_CGAP_Brn67"
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Pred. No. 2.4e-95;
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                                                                                   337;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTP/DTP
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
Plate: LLCM285 row: 1 column: 14
High quality sequence stop: 593.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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                                                                                                         Similarity
                                                                                   Conservative
                                                                                                                                                                                   /tissue_type="melanotic melanoma"
/lab_host="DH108 (phage-resistant)"
/note="organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
a 166 c 183 g 105 t
                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="IMAGE:3617749"
/clone_lib="NIH_MGC_20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
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                                                                                                       91.2%;
99.4%;
                                                                                                       Score 336.4; DB 1 Pred. No. 8.2e-94;
                                                                                     Mismatches
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REFERENCE
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Best Local Similarity 97.5

Matches 356; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
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1 (bases 1 to 704)
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National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        quality sequence stop:
                                                                                                                                  /note="Organ: pooled brain, lung, testis; vector: pCMV-SORT6; Site_1: Not1; Site_2: ECORV (destroyed); RNA source anonymous pool of 6 male brains, age range 23-27; 1 male lung, age 27; and 1 male testis, age 69. Library is oligo-dT primed and directionally cloned (ECORV site is destroyed upon cloning). Average insert size 1.8 kb, insert size range 1-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 021. Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5181585"
/clone_lib="NIH_MGC_115"
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                                                                                                                                                                                                                                                                                                                                                                 /lab_host="DH10B"
                                                          89.7%;
                                   Score 331; DB 13;
Pred. No. 4.4e-92;
0; Mismatches 7
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1 (bases 1 to 705)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
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Tissue Procurement: ATCC
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Clone distribution: MGC clone distribution information can be
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              156
                                                                                                                                                                                                                                                                                                                                                     quality sequence stop: 481
/clone_lib="NH_MCC_43"
/clone_lib="NH_MCC_43"
/tlissue_type="normal pigmented retinal epithelium"
/lab_host="PH10B (phage-resistant)"
/note="Organ: eye; Vector: pOTB7; Site_1: XhoI; Site_2: ECORI; CDNA made by oligo-dT priming. Directionally cloned into ECORI/XhoI sites using the following 5; adaptor: GCCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MCC Library. | "
                                                                                                                                                                                                                                                  /db_xref="taxon:9606"
/clone="IMAGE:4875499"
                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGCAACTTTTTGCGGATTGTTCTTGCTTCCAGGCTTTGCGCTGACAATCAGTGCCTACCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 356;
                                                                                                                                                                                                      CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incytte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                            NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                BE736301.1 GI:10150293
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mRNA sequence.
BE736301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BE736301 850 bp mRNA linear EST 15-SEP-2000 601307087F1 NIH_MGC_39 Homo sapiens cDNA clone IMAGE:3641660 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                             http://image.llnl.gov
Plate: LLCM347 row: m column:
                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                           Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                               High quality sequence stop: 662.
                                                                                                                                                                                                                                                                                                                                                                          (bases 1 to 850)
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/tissue_type="adenocarcinoma"
/lab_host="0H10B (phage-resistant)"
/note="Organ: pancreas; Vector: pOTB7; Site_1: XhoI;
/note="Organ: pancreas; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
                                                                     /db_xref="taxon:9606"
/clone="IMAGE:3641660"
/clone_lib="NIH_MGC_39"
                                                                                                                  /organism="Homo sapiens"
                                                                                                                                              Location/Qualifiers
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96.5%;
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                                                                                                                                                                                                                                                                                                                                                                                              340 bp
y183b07.rl Soares infant brain 1
IMAGE:44619 5', mRNA sequence.
H06756
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                                                                                                                          Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. I Tel: 314 286 1800 Fax: 314 286 1810
                                                                                                                                                                                        Contact: Wilson RK
                                                                                                                                                                                                   Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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95.1%;
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following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).

0; Score 321.6; DB 12; Pred. No. 4.3e-89; Mismatches 16; Indels Length 850; 2: Gaps

2

1 GGCAACTTTTTGCGGATTGTTCTTGCTTNCAGGCTTTGCGCTGCAAATCCAGTGCTACCA 60

253 GGCAACTTTTGGCGGATGGTTCTTGCTTCCAGGCTTTGCGCTGCAAATCCAGTGCTACCA 312

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GTGTGAAGAATTCCAGCTGAACAACGACTGGTCCTCCCCCGAGTTCATGGTGAATTGGCA 372 GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCCCCCGAGTTCATTGTGAATT-GCA 119

CGGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGT 179

CGGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGT 432

ACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCT 239 492

ACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCT

TACTGTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAA-AACCCTCTTTGTA TCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTA 299

ACGGGCCAAGGCCCAAGAAAAGGCGAAGTTCTGGCTCGGGCCTCAGGCCCAGGGCTCCGCA ACGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCA 359

611

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

INIB Homo sapiens cDNA clone

mRNA

linear

EST 21-JUN-1995

Mammalia; Eutheria; Primates; Catarrhini; Homin 1 (bases 1 to 340) Hillier, L., Clark, N., Dubuque, T., Elliston, K.,

,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.
The Washl-Merck EST Project
Unpublished (1995)

Louis, MO

Email: est@watson.wustl.edu Insert Size: 1737 High quality sequence stops: 312

Source: IMAGE Consortium, LLNL;
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Insert Length: 1737 Std Error: 0.00
Seq primer: M13RP1 further information contact the

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REFERENCE
AUTHORS
                                                                                                                                               SOURCE
ORGANISM
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BE746601
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTT 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTCCAGCTGAACAACGACTGCTCCTCCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTT 130
                                                                                                                                                                                                                                                                                                                                                          GGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAACGGGCCAAGG
                                                                                                                                                                                                                                                                                                        -CCAAGAAAAGGGGAAGTTCTGCCTTGGCCCTCAGGC 336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGCGGATTGTTCTTGCTTCCAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          333;
                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                   Contact: Robert Strausberg, Ph.D.
                                                                  NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 654)
                                                                                                                                               Homo sapiens
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BE746601.1 GI:10160593
EST.
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                                                                                                                                                                                                            mRNA sequence.
                                                                                                                                                             human
cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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Location/Qualifiers
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/lab_host="DH10B (ampicillin rea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="Soares infant brain lNIB"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:9606"
/clone="IMAGE:44619"
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/db_xref="GDB:417160"
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98.8%;
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Pred. No. 2.7e-89;
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cDNA clone
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Search completed: November Job time: 501.345 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA 119
                                                                                                                                                                          CGGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGT
                                                                                   ACGGGCCAAGGCCCAAGACAAGGGGGAAGTTCTGCCTCGGCC
                                                                                                        ACGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCC 340
                                                                                                                                                                                                                    TCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTA 299
                                                                                                                                                                                                                                                                  CGGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGT
                                                                                                                                                                                                                                                                                                                                                                                                                                               AGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.
Plate: LLCM/50 row: 1 column: 21
High quality sequence stop: 651.
Location/Qualifiers
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/clone_ilb="NIME_MGC_9"
/clone_ilb="NIME_MGC_9"
/tissue_type="sdenocarcinoma cell line"
/tissue_type="sdenocarcinoma cell line"
/tissue_type="sdenocarcinoma cell line"
/tissue_type="sdenocarcinoma cell line"
/tissue_type="sdenocarcinoma cell line"
/lab_host="sdenocarcinoma cell line"
/note="Organ: ovary: Vector: poTB7; Site_1: xhoI; Site_2:
clored: note="sdenocarcinoma; Directionally
clored: note="sdenocarcinoma; berected by oligo-driven cell
clored: GCCACGAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDM synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
21 a 219 c 187 g 126 t 1 others
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/db_xref="taxon:9606"
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     nucleic search, using sw model
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Match
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Gapop 60.0 , Gapext 60.0
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/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2001B.DAT:*
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ABL40348
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ABG54231
AAF22400
AAF64188
AAF93845
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Ovarian carcinoma
Human ovarian anti
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Human secreted pro
Human cDNA encodin
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Ovarian carcinoma
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	777.8 627.8 577.8 577.8 577.8 577.4 449.1 149.6 6.0 149.6
50 51 263 377 1539 256 257 377 377 377 278 288 288 288 288 288	1619 1619 1619 1619 1524 587 1362 2528 444 473 373 374 374 390 390 390
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AALJ3986 AALJ3986 AALJ3987 ABA18284 AAL10167 ABV33163 ABV23163 ABV53839 ABV598332 AAI17949 ABU59482 AAI187949 ABL59481 AAH468474 AAL109577 ABV050770 AAH68974	ABT03277 ABT03281 ABL40345 ABL40349 ABK33543 AAR94186 ABL703279 ABL60347 AAAH50766 ABL78538 ABL79431 ABLB1262 ABL85916 ABL79431 ABLB5916 ABL79854 ABL79854 ABL79859
<u> </u>	Human ovarian carc Ovarian carcinoma Ovarian carcinoma Ovarian carcinoma CDNA encoding huma Primer specific fo Human ovarian carc Ovarian carcinoma Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc Human ovarian canc

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## ALIGNMENTS

RESULT 1
ABT03280
ID ABTC

ABT03280 standard; cDNA; 625 BP.

Human ovarian carcinoma associated coding sequence SEQ ID NO: 210.

05-SEP-2002 (first entry)

ABT03280;

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WPI;
                                                        14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
                                                                                                                              Homo sapiens.
                                                                                                                                           Human; ovarian cancer; ovarian carcinoma; gene therapy; immunotherapy; cytostatic; gene; ss.
                            Xu J,
                                                                                    13-NOV-2001; 2001WO-US45395
                                                                                                  23-MAY-2002
                                                                                                                WO200239885-A2
                                         (CORI-) CORIXA CORP.
              2002-500186/53.
                          Stolk JA, Algate
                            PA,
                           Fling SP,
                            Molesh DA;
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Novel ovarian cancer polypeptide and polynucleotide, useful for

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RESULT 2
ABL40348
ID ABL4
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AC ABL4
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AC ABL4
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer
 Ovarian cancer;
                          Ovarian carcinoma O1034C EST clone nucleotide sequence
                                                       28-JUN-2002
                                                                                                          ABL40348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2;
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                                                                                                                                                                         TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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                                                                                                                                                                                                                                                                                                            CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
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                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 625 BP; 182 A; 144 C; 182 G; 116 T; 1 other;
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                                                     (first entry)
cancer therapy; vaccine; gene therapy; tumour;
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No. 5.2e-239;
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                                                                                                                                                                                                                                                                                                                         that may be utilised in cancer therapy, for example in a vaccine or complete therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or capanding T-cells specific for a tumour protein, and for inhibiting the expanding T-cells specific for a tumour protein, and for inhibiting the cancer in a patient. They are also useful for complete the complete that and for determining the presence of a cancer in a patient.

The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and complete the complete the complete that the complete the complete that the preparing ribozyme molecules for inhibiting expression of tumour cells. Polypeptides and polynucleotides of the convention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma ol034C EST clone nucleotide
                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 625; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
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(STOL/)
(ALGA/)
                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                 sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ovarian tumour polypeptide and polynucleotide useful in di
prevention and/or treatment of cancer, especially ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
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 GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
             GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
                                                    CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                                                  CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAAGGGGAG
                                                                                                        CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC
                                                                                                                                                           GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                        GCCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
                                                                                                                                                                                                                AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
                                                                                                                                                                                                                              AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
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) STOLK J A.
) ALGATE P A.
) FLING S P.
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2000US-0640173.
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Pred. No. 5.2e-239;
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The present invention describes a composition (I) comprising: carriers and immunostimulants; and a polypeptide (II) of a ovarian tumour polypeptide encoded by a polynucleotide (III) having a cDNA sequence (S1) from the 10912 nucleotide sequences as given in ABL77023 to ABL87934, (III) encoding (II) having a sequence (S2), a T cell population of (II), or antigen presenting cells that express (II). (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to (S1) can be used for detecting ovarian cancer in a patient's biological sample preferably serum or ovarian tissue. The method comprises contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to patient with comparing the amount to
                                                                                                                                                                           Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide
                                                                                                                                                                                                                                     WPI;
                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                                                           Algate PA,
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                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                Human ovarian cancer related DNA clone SEQ ID NO:10876.
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                                                                                                                                                     ID 10876; 489pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                          cancer;
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AC ABT0
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DE Huma
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KW Huma
KW Cyto

05-SEP-2002

(first entry)

ABT03284 standard;

CDNA; 1897

cytostatic;

gene;

Human; ovarian cancer;

Human ovarian carcinoma associated coding sequence SEQ

ovarian carcinoma; gene therapy; immunotherapy;

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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           patient, where the amount of polynucleotide hybridising to (IV) is detected preferably by polymerase chain reaction (PCR). (I) comprising (III) and/or (II) is useful for stimulating and/or expanding T cells specific for an ovarian tumour protein comprising contacting T cells with (III) or (II). (III) is useful in design and preparation of ribozyme molecules for inhibiting expression of the tumour polypeptides and proteins in tumour cells; and to isolate a full length gene from a suitable library e.g., a tumour cDNA library using well known
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                                                               CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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ACAGTGNAAAAAAAAAAAAAAAAA 625
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Pred. No. 5.2e-239;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-NOV-2000; 2000US-0713550.
03-APR-2001; 2001US-0825294.
02-OCT-2001; 2001US-0970966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1897 BP; 435 A; 521 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2; Page 196; 197pp;
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                                                TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
                                                                                               TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG
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RESULT 5
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                that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are cuseful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the certain and incomment of a cancer in a patient. They are also useful for stimulating an immune response in a patient. They are also useful for cancer in a patient and for determining the presence of a cancer in a patient. They isolated polynucleotides of the invention are useful for their capitity to selectively form duplex molecules with complementary stretches confirm the entire desired gene or gene fragments, and for designing and core paring ribozyme molecules for inhibiting expression of tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current
                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                        (XUJJ/)
(STOL/)
(ALGA/)
                                                                                                                                                                                                                                                                                                                                      WPI;
           sequence
                                                                                                                                                                                                                       The invention relates to ovarian tumour polynucleotides and polypeptides
                                                                                                                                                                                                                                                   Claim 1a; Page 127-128; 131pp; English.
                                                                                                                                                                                                                                                                            Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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           represents the ovarian carcinoma 01034C/0591S consensus
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2000US-0640173.
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/product= "Ovarian carcinoma protein Ol034C/O591S"
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                                 Human; ovarian antigen; ovary; ovarian; breast; cancer; tumour; ovarian cancer; breast cancer; tumour; reproductive system disorder; infertility; pregnancy disorder; anovulation; polycystic ovary syndro pcOS; ovarian cyst; dysmenorrhoea; endocrine disorder; infection; inflammatory condition; immune disorder; blood disorder; cardiovascular disorder; respiratory disorder; neurological disorder; gastrointestinal disorder; urinary system disorder; drug screening; gene therapy; chromosome mapping; forensic analysis;
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       gene therapy; chromosome mapping; antibody preparation; cytostatic; antiinflammatory; gynaecological;
                                                                                                                                                            Human ovarian antigen HVCAG04 cDNA,
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            reproductive; chromosome
                      immunomodulatory; neuroprotective,
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AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG

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Similarity

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CAGCGACAGGCGCAGCACCAGCACCTGCACGAACCCCGCCGAAACTGCTGCGAGGACAC

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CC ABP4328) and to cDNAs encoding them (ABC54131-ABC6305), and also CC encompasses polypeptides 90% identical and polynucleotides 95% identical to the sequences of the invention. The invention additionally relates to CC recombinant vectors and host cells comprising human ovarian antigen CC polynucleotides, antibodies against human ovarian antigen and the use C fovarian antigen polynucleotides and polypeptides in diagnosing, CC treating, prognosing or preventing various ovary and/or breast cancer, and CC disorders. Such conditions include ovarian cancer and breast cancer, and CC disorders (e.g., infertility, disorders of pregnancy, anovulation, polycystic ovary syndrome, ovarian cysts, and dysmenorrhoea), endocrine CC disorders, infections (e.g., chamydia, HIV, toxoplasmosis, and toxic vaginitis), immune disorders (e.g., congenital and acquired immunodefictencies, autoimmune cophoritis, systemic lupus erythematosus), blood-related disorders (e.g., anaemia), cardiovascular disorders, candurated ovarian antigen cophoritis, systemic lupus erythematosus), compunitate ovarian antigen expression or activity. The polynucleotides may also be used in screening for compounds which conjugated in disease diagnosis, drug targeting and phenotyping. The present sequence represents cDNA encoding a human ovarian antigen of the conjugated conjugated constitutes and conjugated conjugated conjugated constitutes and conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated conjugated c
                                              specification, but was obtained in electroni
at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated nucleic acid molecules encoding novel ovarian polypeptides, useful in the prevention, treatment and diagnosis of cancer (e.g. ovarian cancer), immune disorders, cardiovascular disorders and
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Sequence 1608 BP; 402 A; 402 C; 417 G; 381 T; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to 2175 novel human ovarian antige ABP43228) and to cDNAs encoding them (ABQ54131-ABQ56395)
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                                                                             sequence data for this patent did not form part of the printed tion, but was obtained in electronic format directly from WIPO
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other;
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Isolated nucleic acid molecule encoding a human secreted protein used in preventing, treating or ameliorating a medical condition  $% \left( 1\right) =\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +$ 

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AAF22400/c
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            P-PSDB;
                       WPI; 2000-647420/62
                                                                                                 09-APR-1999; 99US-0128694
20-JAN-2000; 2000US-0176931
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                                            Ruben SM,
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                                                                                                                                                                                           Human secreted proteins AAB75506 - AAB75554 are encoded by polynucleotide sequences AAF64176 - AAF64224. The specification includes amino acid sequences AAB75555 - AAB75606 which represent fragments of the human secreted proteins, and protein sequences with which they share homology. The proteins and polynucleotides, their agonists and antagonists have activities dependent on the tissues and cells in which they are expressed, examples of these activities include, immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; varotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; polynucleotides, agonists and antagonists can be used to treat or detect
or diagnose various diseases and disorders including, autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. Alzheimer's disease, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin ageing due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to
                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid molecules encoding human secreted proteins, used preventing, treating or ameliorating a disorder, e.g. Alzheime Parkinson's diseases and cancers -
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                                               rheumatoid arthri
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11-JAN-2000;
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                  CCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCA
                                                                                         CATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGAT
                                                                                                                                                                 CAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGAT 246
                                                                                                                                                                                                   CAGGCGGCAGCACCACCACCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTA
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Isogai T,
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2000JP-0183766.
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7.1e-209;
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rug development –
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                                                                                                                                                                 Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1710
                                                                                The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
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                                                Sequence 1619 BP;
                                                                                                                                  Claim 2;
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                                                                        the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  ovarian cancer; ovarian carcinoma; gene therapy;
                                                                                                                                  Page 189-190; 197pp; English.
                                                                                                                                                                                                                                                                                           ; 2000US-0713550.
; 2001US-0825294.
; 2001US-0970966.
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                                                399 A; 415 C; 432 G; 373 T; 0 other;
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Score 486; DB 24;
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02-OCT-2001; 2001US-0970966
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   cancer polypeptide and polynucleotide, useful for presence of ovarian cancer in a patient, and in
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               Ovarian cancer; cancer therapy; vaccine;
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Query Match
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Matches 606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1619 BP; 399 A; 415 C; 432 G; 373 T; 0 other;
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Pred. No. 7.8e-184;
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(first entry)

gene therapy; tumour;

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15-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                                                                                                that may be utilised in cancer therapy for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current
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(STOL/)
(ALGA/)
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                                           GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
      GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTTGGGTTTGGCCGCAG
              GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAG
                                                                                   GCCCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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) STOLK J A.
) ALGATE P A.
) FLING S P.
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t of
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                   polynucleotide useful in diagnosis, cancer, especially ovarian cancer
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gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient.

cancer

The invention relates

may be utilised

relates to ovarian tumour polynucleotides and polypeptides tilised in cancer therapy, for example in a vaccine or Polypeptides and polynucleotides of the invention are

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their
                                                                                                                                                                                                                  1592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                1352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1112
                                                                            08-MAY-2002
 breast cancer; prostate tumour; rectal tumour; liver tumour;
                                            cDNA encoding human PRO protein, Seq ID No
                                                                                                                                    ABK33543 standard;
               Human; secreted protein; PRO; tumour; lung cancer;
                                                                                                                                                                                                                                              600
                                                                                                                                                                                                                                                                                                         540
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 AGTTCTCCTTGCAGAGGACTGGCGCGCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTTGGCCGCAG 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGA 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CAGCGACAGGCGGCACACACC-TGCACGAACACCCGCGGAAACTGCTGCGAGGACA 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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                                                                                                                                                                                                                                             GACAGTG
                                                                                                                                                                                                                                                                            GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA 479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTGGGCAGACACGAGGAGGGGAGAGACAGAGAAAAAGAAAACACAGCATGAGAAACACA 419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC 1351
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                                                                                                                                                                                                                                                                                                                                      GTAAATGAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA 1471
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTTGGCCGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              606;
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                                                                          (first entry)
                                                                                                                                                                                                                                              606
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99.8%;
                                                                                                                                       CDNA; 1524
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7.8e-184;
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              colon cancer;
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1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCCGCCACAAAGCG 60

Matches

Similarity

100.0%; 62.9%;

Score 393; DB 24; Pred. No. 7e-147; Mismatches

Conservative

0,

0

Indels

0;

0

Sequence

1524

BP; 321 A; 433 C; 435 G; 335 T; 0 other;

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25-JUL-2000;
25-JUL-2000;
26-JUL-2000;
28-JUL-2000;
28-JUL-2000;
23-AUG-2000;
24-AUG-2000;
11-SEP-2000;
11-NOV-2000;
26-NOV-2000;
01-DEC-2000;
20-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-JUL-2000;
25-JUL-2000;
25-JUL-2000;
25-JUL-2000;
                      The PRO polypeptides are useful for diagnosing tumours, especially lung cancer, colon cancer, breast tumour, prostate tumour, rectal tumour or liver tumour. The PRO polypeptides are useful for stimulating the proliferation of, or gene expression, in pericyte cells, for stimulating the proliferation or differentiation of chondrocyte cells, for stimulating the release of tumour necrosis factor-alpha from human blood, for stimulating or inhibiting the proliferation of normal human dermal fibroblast cells. The PRO polypeptide may also be used as molecular weight markers and for tissue typing. The PRO nucleic acids have applications in molecular biology, including use as hybridisation probes, and in chromosome and gene mapping. ABK33536-ABK3367 represent human PRO protein coding sequences of the invention.
                                                                                                                                                                                                           The invention relates to one hundred and twenty two nucleic acids encoding PRO polypeptides. The sequences of the 122 PRO polypeptides, encode human secreted proteins. The PRO nucleic acids, polypeptides, agonists and antagonists are useful for treating a PRO related disorder
                                                                                                                                                                                                                                                                                                                     One hundred and twenty two nucleic acids encoding PRO polypeptides, useful for treating a PRO related disorder and for diagnosing tumours such as lung cancer, colon cancer, breast tumour, prostate tumour, retumour or liver tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Grimaldi JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               pericyte cell proliferation; chondrocyte tumour necrosis factor-alpha; gene; ss.
                                                                                                                                                                                                                                                                                        Claim
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25-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                   P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Baker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GETH ) GENENTECH INC.
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DB; AAU83599.
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2000WO-US23328
2000WS-000000P
2000WO-US30873
2000WS-US32678
2000WS-US32678
2000WS-U747259
2000WS-U734956
2001WS-US34956
2001WS-US34956
2001WS-US34956
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cell proliferation;
                                                                                                                                                                                                                                                                                                                                     prostate tumour, rectal
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AAF94186/c
                                                                                                                                                                                                                                                                                                                                               08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
which encode human secretory or membrane proteins represented by AAB88317 - AAB88419. Included in the invention are primers AAF93917 - AAF94295 and AAAF62232 - AAF62235 which are used to isolate the cDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and diagnosis of diseases associated with inappropriate secretory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1492
                                                                                                                                          This invention relates to nucleic acid sequences AAF93744 - AAF93916
                                                                                                                                                                         Claim 5; SEQ ID 620; 609pp + CD ROM; English.
                                                                                                                                                                                                                      Nucleic acids encoding secretory proteins/membrane proteins, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                           10-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; secretory protein; membrane protein; vac rheumatoid arthritis; diabetes; PCR primer; ss.
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2000JP-0183766
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                                                                                                                                                                                                                                                                                      Nishikawa
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                                                                                                                                                                                                                                                                                   Kawai Y,
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                                                                                                                                                                                                                                                                                   Sugiyama
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (e.g. polymerase chain reactions (PCR)) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and activity. The antibodies and antagonists may also be used as therapeutic agents to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of the polypeptides in samples (e.g. by enzyme linked immunosorbant assay (ELISA). Examples of diseases which may be treated include rheumatoid arthritis and diabetes.
                                                                                                                                                                     cytostatic;
                                                                                                                                                                                  Human; ovarian
                                                                                                         WO200239885-A2
                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                  Human ovarian carcinoma associated coding sequence SEQ ID NO:
                                                                                                                                                                                                                                                 05-SEP-2002
                                                                                                                                                                                                                                                                                                              ABT03279 standard; cDNA; 1362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 587 BP; 117 A; 162 C; 140 G; 162 T; 6 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein/membrane protein expression. The nucleic acids and complementary sequences may also be used as DNA probes in diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                    551
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATG 358
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATT 118
                                                                                                                                                                     gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                      ne; ss.
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                                                                                                                                                                                     ovarian carcinoma;
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 365;
                                                                                                                                                                                                                                                                                                              ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 22;
                                                                                                                                                                                   gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>..</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                   immunotherapy;
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14-NOV-2000; 03-APR-2001;

13-NOV-2001; 2001WO-US45395

23-MAY-2002

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RESULT 17
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      US2002004491-A1
                                                                             Ovarian
                                                                                           Ovarian
                                                                                                         28-JUN-2002
                                                                                                                        ABL40347;
                                                                                                                                                                                                                                                                                                                                                                              Sequence 1362 BP; 263 A; 435
                                                                                                                                                                                                                                                                                                                                                                                              e
E
                                                                                                                                                                                                                                                                                                                                                                                                 The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-OCT-2001; 2001US-0970966
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                                                                                                                                                                                                                                                                                                                            1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                      sapiens
                                                                                                                                                                                                          GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
                                                                                                                                                                                                                               CAGCGACAGGCGGCACCACCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC
                                                                                                                                                                                                                                                                                                GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                                                   CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                                                                                                                                                                                                                                                                        GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                                                                                                                                                                    AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGGCGCTGCACAAAGCG
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                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                             invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stolk JA, Algate
                                                                             cancer;
                                                                                          carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                                                                                                               Page 193; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                        (first entry)
                                                                          cancer therapy; vaccine; gene therapy; tumour; cancer;
                 /*tag= a
/product=
                                Location/Qualifiers
1..1362
                                                                                          GPR39 cDNA
                                                                                                                                                                                                                                                                                                                                                       57.4%;
100.0%;
                                                                                                                                     1362
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                 "Ovarian carcinoma
                                                                                                                                                                                                                                                                                                                                                 0;
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                                                                                                                                                                                                                                                                                                                                                       Score 359; I
Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                              362 G;
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                                                                                                                                                                                                                                                                                                                                                                              302 T; 0
                                                                                                                                                                                                                                                                                                                                                               DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Molesh
                                                                                                                                                                                                                                                                                                                                                       .3e-133;
                  protein GPR39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DA
                                                                                                                                                                                                                                                                                                                                                              Length 1362;
                                                                                                                                                                                                                                                                                                                                                 Indels
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                        301
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Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for a patient and for determining the presence of a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polypuncteotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma compa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (XUJJ/)
(STOL/)
(ALGA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1362 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           that may be utilised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ovarian tumour polypeptide and prevention and/or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Xu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1a;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (FLIN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCGAACGGGCGCTGCACAAAGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention relates to ovarian tumour
CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC
                                                                                       CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                                                                                                                                                                                                                                                                                    CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGAGACAC
                                                                                                                                                                                                                                                                                                                                                                                              GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGTTCTCCTTGCAGAGGACTGGCGCGGGGCGCGAAGAGCGAACGGGCGCTGCACAAAGCG 1156
                                                                                                                                     GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
                                                                                                                                                                                   CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                                                                                                                                                                                                                                                                                                                                                        GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                                                                                                                               CAGCGACAGGCGGCAGCACCAGCACCTGCACGAACCCGGCGAAACTGCTGCGAGGACAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                359;
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) STOLK J A.
) ALGATE P A
) FLING S P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stolk JA, Algate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABB09416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 123-124; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           57.4%;
llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000US-0561778.
2000US-0640173.
2000US-0656668.
2000US-0713550.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001US-0825294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   relates to ovarian tumour polynucleotides and polytilised in cancer therapy, for example in a vaccing Polypeptides and polynucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       263 A; 435 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 359; DB 24;
Pred. No. 2.3e-133;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polynucleotide useful in diagnosis, cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       362 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       302 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 1362;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                           359
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                                                                                         916
                                                                                                                                     300
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RESULT 18
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                                                                Matches
                                                                                                                                                        The present invention relates to a method for identifying a nucleic acid molecule (G protein coupled receptor gene, GPCR 4941) associated with a cardiovascular or tumourigenic disorder. The method comprising contacting a sample containing a nucleic acid molecule with a hybridisation probe or amplification primers and detecting the presence. The invention is used in gene therapy. The method of the invention is used for identifying nucleic acids or polypeptides associated with a cardiovascular or tumourigenic disorder such as aberrant angiogenesis, aberrant vascularisation, artherosclerosis, or ovarian cancer, ischaemia/ reperfusion injury, hypertension, restenosis, arterial inflammation, endothelial cell disorders, diabetic retinopathy, psoriasis, myocardial infarction, Grave's disease and leukaemia. The methods can also detect mRNA or genomic DNA in a sample. The present sequence is G protein coupled receptor (GPCR) 4941 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene therapy; aberrant vascularisation; artherosclerosis; ovarian cancer; ischaemia/reperfusion injury; hypertension; arterial inflammation; psoriasis; endothelial cell disorder; diabetic retinopathy; restensis; myocardial infarction; Grave's disease; G protein coupled receptor; GPCR 4941; vasotropic; hypotensive; antiinflammatory; cytostatic; antidiabetic; antipsoriatic; leukaemia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDS
   1256
                                                                                                                                                                                                                                                                                                                                                                                                                                          cardiovascular and tumorigenic disorders, protein coupled receptor (GPCR)-4941 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD18690 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying nucleic acids for the diagnosis and treatment of cardiovascular and tumorigenic disorders, comprises identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-611743/70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-APR-2001; 2001WO-US13788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Fig 1; 118pp; English.
                                                                                                                               Sequence
                                 -
               AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 1197
                                                                  359;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cardiovascular; tumourigenic disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein coupled receptor (GPCR) 4941 cDNA
                                                                                 Similarity
                                                                                                                                 2528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-199908P.
2000US-0635521.
                                                                                                                             BP; 516 A; 766 C; 677 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers 42..1403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product= "Human GPCR 4941"
/note= "The CDS is specifically claimed in calim 1 of
the specification"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                               57.4%;
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                                                                  0;
                                                                               Score 359;
Pred. No.
                                                                  Mismatches
                                                                                                                                 567 T; 2 other;
                                                                                 2.1e-133;
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204 CCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGAGGATCATGTACGCCCGGAAGT 263

Query Match Best Local Matches

Similarity

56.3%; 99.8%;

Score 352; DB Pred. No. 1.6e 0; Mismatches

DB 22; .6e-130;

Length 444; Indels

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Gaps

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Conservative

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RESULT 19
AAH50766/c
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                                                                                                                                                                                                                                                                                                                                                                                                            Human; cancer specific gene expression; gene age related differential expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH50766 standard; cDNA; 444
                                                            patient.
                                                                            are
                                                                                        The
                                                                                                                                      Novel nucleic acid molecules differentially expressed in brain cancers, useful for ascertaining propensity of cell for malignant phenotype or ascertaining suitability of anti-neoplastic drug candidate -
                                                                                                                                                                                                                  Kroes RA,
                                                                                                                                                                                                                                          (NYXI-) NYXIS NEURO THERAPIES
                                                                                                                                                                                                                                                                    17-NOV-1999;
17-NOV-1999;
                                                                                                                                                                                                                                                                                                          17-NOV-2000; 2000WO-US31809
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Sequence
                          haemangioma, glioblastoma, schwannoma, osteoma and pinealoma. The present sequence is a cancer-associated cDNA of the invention.
                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                    25-MAY-2001
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                                              present invention provides the sequences of 184 cDNA fragments which differentially expressed in cancer cell depending on the age of the ient. They can be used to diagnose and identify treatments for cers, particularly brain cancers such as haemangioblastoma, teratoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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                         sequence is a cancer-associated
 444
                                                                                                              Page 50; 82pp; English.
                                                                                                                                                                                                                    Moskal JR,
BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  associated cDNA #95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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 95
A; 113 C;
                                                                                                                                                                                                                     Yamamoto H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
 94 G; 142 T; 0 other;
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RESULT 20
ABL78538/c
ID ABL7855
XX ABL785
AC ABL785
XX 17-MAY
XX Human;
XX Homo s
XX W02001
XX W02001
XX W02001
XX O6-DEC
XX 25-MAY
XX PF 29-MAY
XX PF 29-MAY
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The present invention describes a composition (I) comprising: carriers and immunostimulants; and a polypoptide (III) of a ovarian tumour polypoptide encoded by a polypuctide (III) having a cDNA sequence (S1) from the 10912 nucleotide sequences as given in ABL77023 to ABL87934, (III) encoding (II) having a sequence (S2), a T cell population of (II), or antigen presenting cells that express (II). (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to (S1) can be used for detecting ovarian cancer in a patient's biological sample preferably serum or ovarian tissue. The method comprises contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to
                                                                                                                                                                                                                                                                                                                                                                                                          polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide -
                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Composition for therapy and diagnosis of ovarian cancer comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Algate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ovarian cancer related cDNA clone SEQ ID NO:1516.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Harlocker
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RESULT 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a predetermined cutoff value and thereby detecting ovarian cancer in the patient, where the amount of polynucleotide hybridising to (IV) is detected preferably by polymerase chain reaction (PCR). (I) comprising (III) and/or (II) is useful for stimulating and/or expanding T cells specific for an ovarian tumour protein comprising contacting T cells with (III) or (II). (III) is useful in design and preparation of ribozyme molecules for inhibiting expression of the tumour polypeptides and proteins in tumour cells; and to isolate a full length gene from a suitable library e.g., a tumour cDNA library using well known
                      Composition
                                              WPI; 2002-122075/16.
                                                                                                                                                                                                                                                    Human ovarian cancer related cDNA clone SEQ ID NO:2409
                                                                                                                                                                                                                                                                                                    ABL79431;
                                                                                                                                                                                                                                                                                                                         ABL79431 standard; cDNA;
                                                                                                               26-MAY-2000; 2000US-207484P
                                                                                                                                      29-MAY-2001; 2001WO-US17756
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                                                                                          (CORI-) CORIXA CORP
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                                                                    PΑ,
                                                                                                                                                                                                                               ovarian
for therapy and diagnosis of ovarian cancer comprising of a ovarian tumor polypeptide, polynucleotide encoding antibody specific to polypeptide or T cell expressing
                                                                    Harlocker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                             cancer; ovarian tumour; cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                       606
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polypeptide opolypeptide,

polypeptide, polynucleotide encoding to polypeptide or T cell expressing

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RESULT 22
ABL81262/c
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AC ABL812
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DT 17-MAY
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KW Human;
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KW Homo s
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PN WO2001
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population of (II), or antigen presenting cells that express (II).

(I) has cytostatic activity. An oligonucleotide (IV) that hybridises to (S1) can be used for detecting ovarian cancer in a patient's biological sample preferably serum or ovarian tissue. The method comprises contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to a predetermined cutoff value and thereby detecting ovarian cancer in the patient, where the amount of polynucleotide hybridising to (IV) is detected preferably by polymerase chain reaction (PCR). (I) comprising (III) and/or (II) is useful for stimulating and/or expanding T cells specific for an ovarian tumour protein comprising contacting T cells with (III) or (II). (III) is useful in design and preparation of the tumour polypeptides and proteins in tumour cells; and to isolate a full length gene from a suitable library e.g., a tumour CDNA library using well known
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                                                     Homo sapiens
                                                                                      Human;
                                                                                                                                                          17-MAY-2002
                                                                                                                                                                                           ABL81262
                                                                                                                                                                                                                           ABL81262 standard; cDNA; 409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 349 BP;
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                                                                                                                                                                                                                                                                                                                                              600 GACAGTG 606
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                                                                                                                                                                                                                                                                                                                                                                                  TAACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTAAAAGAAAAGACCATTAAAATGCTTTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA 479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGGGCAGACACGAGCAGGAGGAGGAGACAGAGAAAAAGAAAAACACACGATGAGAAACACACA 419
                                                                                                                                                                                                                                                                                                               GACAGTG 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; SEQ ID 2409; 489pp; English.
                                                                                      ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                          (first entry)
                                                                                                                        cancer related cDNA clone SEQ ID NO:4240
                                                                                      cancer; ovarian tumour; cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         79 A; 84 C;
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                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local :
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nes 313; Conserv
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                                                                            GAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGC
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99.7%;
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Pred. No. 3.6e-95;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                  DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                 Length 409;
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RESULT

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ABL85916/c

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ABL85916 standard; cl
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AC ABL85916;

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DT 17-MAY-2002 (first |
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DT 17-MAY-2002 (first |
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Human ovarian cancer
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Human; ovarian cancer
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Human; ovarian cance
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Human; ovarian cance
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Homo sapiens.

XX

PN 06-DEC-2001.

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29-MAY-2001; 2001W0-
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PN (CORI-) CORIXA CORP.
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Algate PA, Harlocke
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                                                             CAATCAGCCCTGGGCAGACACGAGCAGGAGGAGAGAGAAAA 396
                                                                                                                                                                                                                                                          AGAAGGGGAGGAGCATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTCCTGGGTT 290
CAATCAGCCCTGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAA
                                                                                                                           TGGCCGCAGCCATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAA
                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28
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                                                                                                                                                                                                                                                                                                                                                                                                                               26.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                           Score 166; DB 24; Length 201; pred. No. 1.4e-56;
                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46 G;
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В δÃ 망

207

147

426 GAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTAC

485 206

GAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTAC

366 AGACACGAGCAGGAGGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAGTAAAT 425

Matches Query Match

Similarity

Conservative (

0,

Mismatches

0,

0

Gaps

0

Length 390; Indels

24.6%; Score 154; DB 24; 100.0%; Pred. No. 7.4e-52;

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RESULT 24
ABL78554
                                                           CC ABL87934, (III) encoding (II) having a sequence (S2), a T cell
CC (Population of (II), or antigen presenting cells that express (II).
CC (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to CC (S1) can be used for detecting ovarian cancer in a patient's biological contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to compare the amount of polynucleotide hybridising to (IV) and comparing the amount to patient, where the amount of polynucleotide hybridising to (IV) is detected preferably by polynucrase chain reaction (PCR). (I) comprising (III) and/or (II) is useful for stimulating and/or expanding T cells compared to (IV) is useful for an ovarian tumour protein comprising contacting T cells compared to (III) or (III) is useful in design and preparation of cells contacting in tumour cells; and to isolate a full length gene from a contacting sequence (III) and protein comprising contacting to (IV) is detected proteins in tumour cells; and to isolate a full length gene from a contacting sequence (IV) and compared to (IV) is detected proteins in tumour cells; and to isolate a full length gene from a contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and comparising contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) and contacting sequence (IV) an
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human ovarian
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-MAY-2001; 2001WO-US17756
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present invention describes a composition (I) comprising: carriers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PA, Harlocker
390 BP; 121 A; 82 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID 1532;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer related cDNA clone SEQ ID
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99 G; 86 T; 2 other;
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RESULT 25
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                                                                                                                               Query Match
Best Local Similarity
                                                                                                              Matches
                                                                                                                                                                                                                                      amount of polynucleotide hybridisting to (IV) and comparing the amount to a predetermined cutoff value and thereby detecting ovarian cancer in the patient, where the amount of polynucleotide hybridising to (IV) is detected preferably by polynerase chain reaction (PCR). (I) comprising (III) and/or (II) is useful for stimulating and/or expanding T cells specific for an ovarian tumour protein comprising contacting T cells with (III) or (II). (III) is useful in design and preparation of ribozyme molecules for inhibiting expression of the tumour polypeptides and proteins in tumour cells; and to isolate a full length gene from a suitable library e.g., a tumour CDNA library using well known
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide -
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                                                                                                                                                                                                                                                                                                                                                                                                                                  (S1) can be used for detecting ovarian cancer in a patient's biological sample preferably serum or ovarian tissue. The method comprises contacting a biological sample from a patient with (IV), detecting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABL87934, (III) encoding (II) having a sequence (S2), a T cell population of (II), or antigen presenting cells that express (II).
(I) has cytostatic activity. An oligonucleotide (IV) that hybridises to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200192581-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a composition (I) comprising: carriers and immunostimulants; and a polypeptide (II) of a ovarian tumour polypeptide encoded by a polypucleotide (III) having a cDNA sequence (SI) from the 10912 nucleotide sequences as given in ABL77023 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Algate PA, Harlocker SL,
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                                                                                                                                                                                       Sequence 558
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 198
                                                        138 ACAGCACCTGCACGAACACCCGCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGT 197
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TGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCC 257
                                      ACAGCACCTGCACGAACACCCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGT 193
                                                                                                              Conservative
                                                                                                                                                                                       BP; 147 A; 133 C; 166 G; 106 T; 6 other;
                                                                                                                                 19.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jones
                                                                                                                               Score 120; DB 24; Pred. No. 2.3e-38;
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GGCGCTGTCGGTGGTGCGCATGTACGC

AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 208

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RESULT 26
ABL84848
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                                       Matches
                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide -
                                                                                                                                                        with (III) or (II). (III) is useful in design and preparation of ribozyme molecules for inhibiting expression of the tumour polypeptides and proteins in tumour cells; and to isolate a full length gene from a suitable library e.g., a tumour cDNA library using well known
                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a composition (I) comprising: carrier and immunostimulants; and a polypeptide (II) of a ovarian tumour polypeptide encoded by a polynucleotide (III) having a cDNA sequence (SI) from the 10912 nucleotide sequences as given in ABL77023 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID 7826; 489pp; English.
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                                                                                                          Sequence 381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polypeptide
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                                                                                                                                             techniques.
                                                       Local
   1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
                                                       Similarity
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                                       Conservative
                                                                                                             В₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer; ovarian tumour; cytostatic; gene;
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                                                                                                            A; 101 C;
                                                      14.7%; Score 92;
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             describes a composition (I) comprising: carriers
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                                                                                                             122 G;
                                        Mismatches
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                                                                                                             73 T; 0 other;
                                                                          DB 24;
                                                         3.2e-27;
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GGCGCTGTCGGTGGTGGAGTGCGCATGTACGC

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ID AAL33985
                                                                                                                                             Matches
                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               variants of proteins related to amylases, amyloid proteins, angiopoletin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                        diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; anglopoletin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                      Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to oligonucleotides encoding polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections - \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2} \frac{1}{2
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27-DEC-2000; 2000US-0173419.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunosuppressive; immunostimulatory; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                             leukaemia),
                                                                    293
26
                                                                                                                                                                           Local
                                                                GCCGCAGCCATGATCCTCCGAATCTG 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001-465210/50
                                                                                                                                         Similarity
26; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 3451; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                          BP; 14
                                                                                                                                                                                                                                                                                                                                                                                             diseases of the nervous system and an infection
                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Leach
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                      13 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               50
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                                                                                                                                                                                                                                                                                   17 G; 6 T; 0 other;
                                                                                                                                         0;
                                                                                                                                                                           Score 26;
Pred. No.
                                                                                                                                         Mismatches
                                                                                                                                                                                   ö
                                                                                                                                                                               0.0
BB
                                                                                                                                                                           . 68;
                                                                                                                                         0;
                                                                                                                                                                                                          Length 50;
                                                                                                                                         Indels
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                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                             of pathogenic
                                                                                                                                     Gaps
                                                                                                                                     0
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RESULT 28
AAL33986/c
                                                                                                      Query Match
Best Local S
Matches 26
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                                                                                                                                                                                                                                                                                                                                                                                          apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune
                                                                                                                                                                                                                                                                                      diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
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                                                                                                                                                                                                             Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoletin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-DEC-1999;
27-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nervous system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human SNP oligonucleotide #7194.
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                                                  ACCTCGTCCAGTCGTGCTTGGGTTTG 292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001-465210/50
  ACCTCGTCCAGTCGTGCTTGGGTTTG
                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 3451; 4143pp; English.
                                                                                                                                                                                                                BP;
                                                                                                      Conservative
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                                                                                                                                4.2%;
100.0%;
                                                                                                                                                                                                             12 C; 17 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             autoimmune disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   50
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                                                                                                      0;
                                                                                                                                Score 26;
Pred. No.
                                                                                                      Mismatches
25
                                                                                                                                NO.
                                                                                                                                DB 22;
0.68;
                                                                                                                                                      Length 50
                                                                                                      Indels
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                                                                                                      Gaps
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AAL27277/c ID AAL27277

standard;

DNA;

51

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RESULT

29

В

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ABA18284
ID ABA:
XX
AC ABA:
XX
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                                                                            RESULT 30
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                                                                                                                                                                                  Query Match
Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating fact complement related protein; cytochrome; kinesi; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
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                  ABA18284;
                                                                                                                                                                                                                                            Sequence 51 BP;
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27-DEC-2000; 2000US-0173419
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic;
                                               ABA18284 standard;
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                                                                                                                                      401 AACACAGCATGAGAACACAGTAAAT 425
                                                                                                                        51
                                                                                                                          AACACAGCATGAGAACACAGTAAAT
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                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease; autoimmune disease; infection;
disease; ss.
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                                                                                                                                                                                                 100.0%;
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                                               ВP
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Pred. No.
                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                              other;
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                                                                                                                                                                                                                                                                                       of pathogenic
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Human; nootropic; neuroprotective; cytostatic; dermatological; virucide; immunosuppressive; antiinflammatory; anti-HIV; antibacterial; vulnerary; antiparkinsonian; antistckling; antianaemic; antiarthritic; cancer; antirheumatic; hepatotropic; cerebroprotective; antiinflammatory; antiallergic; antidiabetic; antiuleer; anticonvulsant; antifungal; antiparasitic; cardiant; immune disorder; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
                                                                                                                                                                                                                                                                                                                                                                                             neurological disease; infection; nephrotropic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               system related
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28-JUN-2000 30-JUN-2000 07-JUL-2000 11-JUL-2000 11-JUL-2000 11-JUL-2000 11-JUL-2000 11-JUL-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AUG-2000 11-AU WO200159063-A2 Homo sapiens. 17-JAN-2001; 2001WO-US01334 16-AUG-2001. 2000US-0224518
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2000US-0237039.
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2000US-0237040.
2000US-0239935.
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2000US-0235498
2000US-0235834
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AAL10167/c
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Best Local Similarity
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08-DEC-2000;

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08-DEC-2000;

08-DEC-2000;

01-DEC-2000;

01-DEC-2000;
          10-JAN-2001; 2001WO-US00798
                                       19-JUL-2001
                                                                                                                     Human; breast cancer; cell marker; cytostatic; ss
                                                                                                                                                Human breast cancer expressed polynucleotide 2624.
                                                                  WO200151628-A2
                                                                                             Homo sapiens.
                                                                                                                                                                               07-DEC-2001
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cDNA; 377

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Indels

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The invention relates to novel genes (ABAl1004-ABA21534) and proteins (ABB14678-ABB18001) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune thyroiditis, diabetes mellitus, Crohn's clieses, multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as wiral, bacterial, fungal cardiorassitic infectious diseases such as viral, bacterial, fungal cardiorassitic infectious diseases such as viral, bacterial, fungal cardiorassitic infections.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                              Sequence 263 BP; 88 A; 36 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 10615; 1701pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acids encoding 3224 human nervous system antigen polypeptides, useful for preventing, diagnosing and/or treating nervous system cancers and metastases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-541565/60.
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3.7%;
ilarity 100.0%;
Conservative
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: 2000US-0251479.

2000US-0251469.

: 2000US-0251868.

: 2000US-0251869.

: 2000US-025189.

: 2000US-025199.

: 2000US-025199.

: 2000US-0254097.

: 2001US-0259678.
                             Score 23; DB 22; Length 263; Pred. No. 8.7;
                                                                                                                                                         67 G;
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14-MAR-2000; 2000US-0199167.
24-MAR-2000; 2000US-0193099.
29-MAR-2000; 2000US-0193480.
15-MAY-2000; 2000US-0205230.
09-JUN-2000; 2000US-0213115.
25-JUL-2000; 2000US-0220534.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to human breast cancer expressed polynucleotides (AALO7544-AAL26789) and methods of assessing whether a patient is afflicted with breast cancer by examining the correlation between the expression of certain markers and the cancerous state of breast cells. The polynucleotides and encoded polypeptides are potential markers for the content of the content of the cancerous state of breast cells.
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                                                                                                                                                                                                                                                                                                                 pharmacogenomic
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                                                                                                                                                                       17-FEB-2000;
                                                                                                                                                                                                    20-FEB-2001;
                                                                                                                                                                                                                                                                                                                                   Human;
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                                        Schlegel R,
                                                                    (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                 18-JUL-2000;
                                                                                                                             09-JUN-2000;
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2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-211314P.
2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                        Endege
                                                                                                                                                                                                    2001WO-US05171
                                                                                                                                                                                                                                                                                                                 cancer; cytostatic; carcinogen; pharmacodyanamic marker;
marker; gene; ss.
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pred. No. 8.
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16-MAR-2000; 2000US-189862P.
25-MAY-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-219007P.
13-DEC-2000; 2000US-255281P.
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(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usel for detecting presence of prostate cancer, stage of prostate cancer
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in a patient;
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                                                                                                         Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usel for detecting presence of prostate cancer, stage of prostate cancer % \left( 1\right) =\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 1\right) +\left( 
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                                                                                                                                                                                                                                                                                                                                                Schlegel R,
                                                                                                                                                                                                                                                                                                                                                                                                                         (MILL-) MILLENNIUM PREDICTIVE MEDICINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        is also useful as a pharmacodyanamic or pharmacogenomic marker:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytostatic; carcinogen; pharmacodyanamic marker;
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Claim

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Matches 23
The invention relates to an isolated polynucleotide (I) comprising sequences selected from 273 sequences fully defined in the specification. (I), including its encoded polypeptide (II), an antibody binding to (II), a fusion protein comprising (II) and a T-cell population stimulated by (I) or (II) are useful for stimulating an immune response in a patient and treating head and neck cancer in a patient. An oligonucleotide (III) that hybridises to (I) is useful for determining the presence of cancer in a patient, by obtaining a biological sample from the patient, contacting the sample with (III), detecting in the sample an amount of a polynucleotide that hybridises to the oligonucleotide, and comparing the
                                                                                                                                                                                                                                                Wang
                                                                                                                                               Claim 1; Page
                                                                                                                                                                                                Novel polynucleotide
                                                                                                                                                                                                                        WPI; 2002-257467/30
                                                                                                                                                                                                                                                                                                  03-AUG-2000;
16-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             gene;
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(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
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                                                                                                                                                                                 pel polynucleotide encoding head and neck tumour pharmaceutical compositions, e.g. vaccines, for
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2000US-249933P
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                                                                                                                                              English.
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selecting a composition for inhibiting prostate cancer i assessing the prostate cell carcinogenic potential of a determining whether prostate cancer has metastasized in assessing the aggressiveness or indolence of prostate cancer than the cancer is a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to be a second to

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                                                                                            The invention relates to an isolated nucleic acid molecule (I) com a nuclectide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;

(a) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
                                                                                                                                                                                                                                                                                                                                                                              17-FEB-2000; 2000US-183319P.
16-MAR-2000; 2000US-189862B-25-MAR'2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-219007P.
                                                                                                                                                                                                               Claim 1;
                                                                                                                                                                                                                                   Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, uses for detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amount of polynucleotide that hybridises to the oligonucleotides to a predetermined cut-off value. (I) and (II) are useful in pharmaceutical compositions, e.g. vaccines, and other compositions for the diagnosis; treatment of head and neck cancer. ABK53806-ABK54078 represent human hand neck cancer cDNA sequences of the invention.
                                                                                                                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABV08332 standard; cDNA; 377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200160860-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABV08332;
                                                                                                                                                                                                                                                                                                                                           (MILL-) MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-FEB-2001; 2001WO-US05171
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                                                                                                                                                                                                             Page 1324; 11750pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic; carcinogen; pharmacodyanamic marker;
                                                                                                                                                                                                                                                                                                                   Monahan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      marker
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Pred. No.
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22;
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                                                                                                                                                                         ) comprising of the
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226 GTGNAAAAAAAAAAAAAAAAAA 247

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RESULT 36
AAI87949
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Best Local
                                            Matches
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                                                                                                                                                                                                                                                                     The invention relates to human polynucleotides (AAI79941-AAI93841) and the encoded proteins (AA000010-AA013910) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiseis regulating activity, tissue growth factor activity, immunomodulatory activity and activinyinhibin activity and may be useful in the diagnosis and/or reperment of cancer lawbeamia across current diagnosis and/or reperment of cancer lawbeamia across current diagnosis and/or reperment of cancer lawbeamia across current diagnosis and/or reperment of cancer lawbeamia across current diagnosis and/or reperment of cancer lawbeamia across current diagnosis and/or reperment of cancer lawbeamia across current diagnosis and/or reperment of cancer lawbeamia across current diagnosis and/or activity when the content of cancer lawbeamia across current diagnosis and/or activity when the content of cancer lawbeamia across current diagnosis and/or activity when the cancer lawbeamia across current diagnosis and/or activity when the cancer lawbeamia across content according to the cancer lawbeamia activity and activity and activity and activity according to the cancer lawbeamia activity and activity according to the cancer lawbeamia activity according to the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia activity and the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia according to the cancer lawbeamia according to the can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; cytokine; cell proliferation; cell differentiation; gene th vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia; nervous system disorders; arthritis; inflammation; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Isolated nucleic acids and polypeptides, useful for preventing diagnosing and treating e.g. leukaemia, inflammation and immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P-PSDB; AA008018.
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18-MAY-2000;
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                                                                                                                                                                                          Note: The sequence data for this patent did not form specification, but was obtained in electronic format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            604 GTGNAAAAAAAAAAAAAAAA 625
                                                                                                                              Sequence
                                                                                                                                                                                                                                         inflammation.
                                                                                                                                                                                                                                                             treatment of cancer,
  604 GTGNAAAAAAAAAAAAAAAAA
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                                                                                                                                                                        ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polynucleotide SEQ ID NO 8009
                                              22; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 8009; 1399pp + Sequence Listing; English.
                                                                                                                                442
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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2000US-0577409.
                                                                                                                              BP;
                                                                                                                                167
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                                                                3.5%;
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                                                                                                                                                                                                                                                             leukaemia,
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                                                                                                                                  65 C;
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Pred. No.
                                                                Score 22;
pred. No.
      625
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                                                                                                                                  102 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                             nervous system disorders, arthritis and
                                              red. No. 20 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cell differentiation; gene therapy;
                                                                                                                                  85 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T; 27
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                                                                                        DB
                                                                    20;
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                                                                                                                                     23 other;
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                                                                                        Length 442;
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                                                Gaps
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                                                                      Matches
                                                                                                       Query Match
                                                                                                                                                                                                                                            The nucleic acids, proteins, antibodies and (antiagonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis, (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as viral, bacterial, fungal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer; vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic; cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL90482 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel 1405 isolated polypeptides, useful for diagnosis, treatment and prevention of neural, immune system, muscular, reproductive, gastrointestinal, pulmonary, cardiovascular, renal and proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neurological disease; infection; human; secreted protein; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human polynucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                         (ABB89040-ABB90444) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-MAY-2000; 2000US-205515P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
Sequence 1329 BP;
                                                                                                                                                                           Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to novel genes (ABL89449-ABL90853) and proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; SEQ ID NO 1044; 2081pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-122018/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (HUMA-) HUMAN GENOME
                                   604 GTGNAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                               parasitic infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABB90073.
                                                                                        Similarity
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                                                                       Conservative
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                                                                                                                                            263 A; 407 C; 369 G;
                                                                                        100.0%;
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                                                                                                            3.5%;
                                                                       s; Score 22; DB; Pred. No. 18; 0; Mismatches
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   1315
                                                                                                                                               288 T; 2 other;
                                                                                                            DB
                                                                                            18;
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                                                                             Indels
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                                                                           Gaps
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cc encoding an Arabidopsis thaliana transcription factor, their variants, complements, fragments, or related polynucleotide with 31% to 95% cc sequence identity, where the plant possesses an altered trait as compared to a wild-type or reference plant, or the plant axhibits an altered cc phenotype as compared to a wild-type or reference plant, or the plant compared to a wild-type or reference plant, or the plant cc associated with a plant trait as compared to a wild plant. Also included created with a plant comprising the polynucleotides, a computer cc areadable medium having stored sequence information, and identifying a compose comprising inputting sequence information, and identifying a sequences comprising inputting sequence information. The isolated or recombinant polynucleotide is used for producing a plant having a modified trait, the method comprising selecting a polynucleotide that cencodes a polypeptide or an antisense nucleic acid, inserting the polynucleotide or antisense nucleic acid, inserting the compant of the polypeptide or antisense nucleic acid, thereby producing a modified trait, and selecting for a modified trait (e:g. increased producing a modified compant of a plant or a cell of a plant to overexpress the polypeptide or antisense nucleic acid, thereby producing a modified compant of agriculturally useful proteins or metabolic chemicals,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 38
ABK65378
                                                                                                                                                                                                                                                                                                                           The invention relates to 1 of 232 isolated or recombinant polynucleotides encoding an Arabidopsis thaliana transcription factor, their variants,
                                                                                                                                                                                                                                                                                                                                                                            Claim 4; Page 927-929; 941pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-292022/33.
P-PSDB; AAU93192.
                                                                                                                                                                                                                                                                                                                                                                                                                          An isolated or recombinant polynucleotide used to produce a transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-AUG-2000; 2000US-227439P.
16-NOV-2000; 2000US-0713994.
16-APR-2001; 2001US-0837944.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               agriculture; metabolic chemical; environmental stress; drought; microbial disease resistance; herbicide resistance; seed yield; fruit yield; growth rate; leaf senescence; flower senescence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pilgrim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200215675-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HEAR/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REUB/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RATCLIFF O. REUBER J L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HEARD J.
JIANG C.
KEDDIE J.
ADAM L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PILGRIM M.
CREELMAN R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 M, Creelman
Ratcliff O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PINEDA O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MENDEL BIOTECHNOLOGY INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RIECHMANN J L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DUBELL A J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene; transcription factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA encoding a transcription factor #230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dubell AJ, Heard J, Juber JL, Riechmann JL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jiang C,
L, Yu G,
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Pineda O;
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RESULT 3
AAH34281
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                                                                                                                                 cancer associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens. The colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer associated Ps, by inserting the nucleic acids into a host cell and culturing the cell for expression and context the colon cancer associated Ps, by inserting the nucleic acids into a host cell and culturing the cell for expression and context associated Ps.
     present invention.

N.B. Pages 666 to 682 and page 7053 of the sequence listing were missing at time of publication, meaning no sequences are present SEQ ID NO:1027 to 1052, 7921 and 7922.
                                                               by inserting the nucleic actus the design the prevention, to express the proteins. N and P can be used in the prevention, and treatment of colorectal carcinomas and cancers. AAH37196 to and AAB77789 represent sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                         AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human
                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids encoding 4277 human colon cancer-associated polypeptides, useful for preventing, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-235357/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-SEP-1999;
03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    specification). The present sequent specification is that it and transcription factor. encoding an A. that it and transcription factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pest tolerance, environmental stress response (e.g. drought), microbial disease resistance, herbicide resistance, seed and fruit yield, growth rate, leaf and flower senescence and many other traits listed in the specification). The present sequence is one of the 232 polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P-PSDB; AAG74876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; colon cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH34281;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     colon
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                                                                                                                                                                                                                                                                                                                                                                                Page 3077-3078; 9803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0157137
99US-0163280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          colon cancer antigen; diagnosis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Birse CE,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rosen CA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB
18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24; Length 1388;
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                                                                                    on, diagnosis to AAH37204 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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for

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RESULT 40
ABL90818/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antifucer; vulnerary; anticonvulsant; antibacterial; antifungal; antibarastic; cardiant; gene therapy; cancer; lumune disorder; cardiovascular disorder; neurological disease; infection; human; secreted protein; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1823 BP; 355 A; 609 C;
                                                                              disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as viral, bacterial, fungal and parasitic infections.
                                                                                                                                                         The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's haemolytic anaemia, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                   Novel 1405 isolated polypeptides, useful for diagnosis, treatment and prevention of neural, immune system, muscular, reproductive, gastrointestinal, pulmonary, cardiovascular, renal and proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                Note: The sequence data for this patent did not form part of printed specification, but was obtained in electronic format
                                                                                                                                                                                                                                                  (ABB89040-ABB90444) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful
                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ ID NO 1380; 2081pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             604 GTGNAAAAAAAAAAAAAAAAA 625
   Sequence
                                                                                                                                                                                                                                                                                                                          invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABB90409.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard; cDNA;
                                 at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rosen
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                                                                                                                                                                                                                                                                                                                          relates
   BP;
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 383 A; 571 C; 686 G;
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                                                                                                                                                                                                                                                                                                                      to novel genes (ABL89449-ABL90853) and proteins
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Pred. No.
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17;
   400 T; 9 other;
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                                                      part of the format directly
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Query Match

22;

DB

Length 2049;

07-DEC-2001

(first entry)

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RESULT 41
AAH69474/c
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RESULT 42
AAL09957/c
ID AAL09
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AC AAL09
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DT 07-DE
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                                                                                                                                             Matches
                                                                                                                                                        Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH69474 standard; cDNA; 259
                                                                                                                                                                                                            cervical cancer with cytostatic activity. The nucleic acids and encoded polypeptides are useful: to assess if a patient is afflicted with cervical cancer or has a pre-malignant condition; to monitor the progression of cervical cancer or a premalignant condition in a patient; and to select and/or assess the efficacy of a compound or therapy for inhibiting cervical cancer in a patient. The nucleic acids may also be useful for gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cervical cancer marker nucleic acid 748
                                                                                                                                                                                                                                                                                                                                     and
                                                                                                                                                                                                                                                                                                                                                 New
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21-JUL-2000;
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12-MAY-2000;
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21-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cervical cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-SEP-2001
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                                                                                                                                                                                          Sequence 259 BP; 73 A; 33 C;
                                                                                                                                                                                                                                                                                          The invention relates to novel genes (AAH68727-AAH73383) associated with
                                                                                                                                                                                                                                                                                                               Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                            Schlegel
                                                                                                                                                                                                                                                                                                                                                                                                                (MILL-) MILLENNIUM PREDICTIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-DEC-2000;
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                        AAL09957;
                                            AAL09957
                                                                                                                       605 TGNAAAAAAAAAAAAAAAA 625
                                                                                                    63
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                                                                                                                                                                                                                                                                                                                                    isolated nucleic acid for diagnosing and treating cervical cancer for assessing and detecting compounds for treating the cancer \boldsymbol{\cdot}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GTGNAAAAAAAAAAAAAAAA 304
                                                                                                                                                                                                                                                                                                                                                                       2001-375006/39
                                                                                                    TGNAAAAAAAAAAAAAAA 43
                                                                                                                                               21;
                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                               Page 232; 1051pp; English.
                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                      2000US-0189315.
2000US-0203791.
2000US-0210600.
2000US-0220114.
                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                            Deeds J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytostatic; pre-malignant condition;
                                             CDNA; 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                          100.0%;
                                                                                                                                                                     3.48;
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                                                                                                                                                         Score 21;
Pred. No.
                                                                                                                                                                                          37 G;
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                                                                                                                                                                                                                                                                                                                                                                                              Zhao
                                                                                                                                                                                           87 T;
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                                                                                                                                                         DB 22; Length 259; 54;
                                                                                                                                                                                            29
                                                                                                                                                                                            other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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ABV05091,
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-JAN-2000; 2000US-0176077.
14-MAR-2000; 2000US-0189167.
24-MAR-2000; 2000US-0193480.
15-MAR-2000; 2000US-0193480.
15-MAY-2000; 2000US-0205230.
09-JUN-20000; 2000US-020534.
                          20-FEB-2001; 2001WO-US05171.
                                                                                                  WO200160860-A2
                                                                                                                                     Homo sapiens
                                                                                                                                                                       Human; prostate cancer;
pharmacogenomic marker;
                                                                                                                                                                                                                         Human prostate expression marker cDNA 5082.
                                                                                                                                                                                                                                                                                                                                     ABV05091 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to human breast cancer expressed polynucleotides (AALO7544-AAL26789) and methods of assessing whether a patient is afflicted with breast cancer by examining the correlation between the expression of certain markers and the cancerous state of breast cells. The polynucleotides and encoded polypeptides are potential markers for detecting, diagnosing, monitoring, characterising treating and potentially preventing breast cancer. The polynucleotides and encoded polypeptides are also useful for isolating compounds with cytostatic
                                                             23-AUG-2001.
                                                                                                                                                                                                                                                                13-SEP-2002
                                                                                                                                                                                                                                                                                                      ABV05091;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 288 BP; 76 A; 53 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 462; 3695pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New peptide useful as a marker for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lillie J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
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                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                       CDNA; 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.4%;
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                                                                                                                                                                                     cytostatic; carcinogen;
                                                                                                                                                                       gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 22;
54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis of breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 288;
                                                                                                                                                                                   pharmacodyanamic marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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ID ABV057
AC XXX DIT CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF CONTROL OF 
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Query Match
Best Local :
    09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                                                                   17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
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16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                            20-FEB-2001;
                                                                                                                                                                                                      23-AUG-2001;
                                                                                                                                                                                                                                                  WO200160860-A2
                                                                                                                                                                                                                                                                                                                                      pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                        Human; prostate cancer; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                Human prostate expression marker cDNA 5761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABV05770;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABV05770
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in a patient;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    is also useful as a pharmacodyanamic or pharmacogenomic marker.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
; 2000US-183319P.
; 2000US-189862P.
; 2000US-207454P.
; 2000US-211314P.
; 2000US-2119007P.
; 2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; 2000US-183319P.
; 2000US-189862P.
; 2000US-207454P.
; 2000US-211314P.
; 2000US-219007P.
; 2000US-255281P.
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                                                                                                                                                         2001WO-US05171.
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                                                                                                                                                                                                                                                                                                                                      gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 21;
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                                                                                                                                                                                                                                                                                                                                                      carcinogen;
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                                                                                                                                                                                                                                                                                                                                                    pharmacodyanamic marker;
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AAH68974/c
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Best Local
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(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer, stage of prostate cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (d) assessing the efficacy of a therapy for inhibiting prostate cancer
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                                                                                                                                                                                                                                                                                                                                                                                       08-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200142467-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cervical cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human cervical cancer marker nucleic acid 248.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH68974 standard; cDNA; 359 BP
                                                                                                                                                                                                                                                14-MAR-2000;
12-MAY-2000;
                                                                                                                                                                                                                                                                                                                                 08-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                              14-JUN-2001.
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                            WPI; 2001-375006/39
                                                                              Schlegel R,
                                                                                                                                     (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                                                                                                                         21-JUL-2000;
                                                                                                                                                                                                                        09-JUN-2000;
                                                                                                                                                                                                                                                                                                      21-DEC-1999;
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les 21; Conserv
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2000US-0203791.
2000US-0210600.
2000US-0220114.
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                                                                                                                                                                                                                                                                                                                                                                                       2000WO-US33312
                                                                                Deeds J,
                                                                                                                                                                                                                                                                                                      99US-0169681.
99US-0171350.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cytostatic; pre-malignant condition; gene therapy; ss
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                                                                                                                                                                                                                                                                            The invention relates to novel genes (AAH68727-AAH73383) associated with cervical cancer with cytostatic activity. The nucleic acids and encoded polypeptides are useful: to assess if a patient is afflicted with cervical cancer or has a pre-malignant condition; to monitor the progression of cervical cancer or a premalignant condition in a patient; and to select and/or assess the efficacy of a compound or therapy for inhibiting cervical cancer in a patient. The nucleic acids may also be needed to be a constant of the constant of the constant of the cancer in a patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated nucleic acid for diagnosing and treating cervical cancer and for assessing and detecting compounds for treating the cancer \cdot
                                                                                                                                                                                                                         Sequence 359 BP; 96 A; 62 C; 48 G;
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CURRENT APPLICATION NUMBER: US/09/149,476
CURRENT FILING DATE: 1998-09-08
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ER APPLICATION NUMBER: 60/047,594

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ER FILING DATE: 1997-04-11
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R APPLICATION NUMBER: 60/056,910
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ER APPLICATION NUMBER: 60/040,163

ER FILING DATE: 1997-03-07

ER APPLICATION NUMBER: 60/047,600

ER FILING DATE: 1997-05-23

ER APPLICATION NUMBER: 60/047,615

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APPLICATION NUMBER: 60/056,884
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FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,903
FILING DATE: 1997-08-22 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/0 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/ FILING DATE: 1997-08-22 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/ FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/ FILING DATE: 1997-08-22 APPLICATION FILING DATE: APPLICATION 1 APPLICATION NUMBER: APPLICATION FILING DATE: FILING DATE: NUMBER: 60 NUMBER: 60: 1997-08-7 NUMBER: 60/0: 1997-08-22 NUMBER: 60/0: 1997-08-22 NUMBER: 60/ : 1997-08-22 NUMBER: 60/0: 1997-08-22 NUMBER: 60/ : 1997-08-22 NUMBER: 60/056,636: 1997-08-22 NUMBER: 60/ : 1997-08-22 NUMBER: 60/ : 1997-04-11 NUMBER: 60/ : 1997-05-23 NUMBER: 60/057,761: 1997-08-22 NUMBER: NUMBER: 1997-08-60/043,576 60 60/056,875 60/056,881 60/056,876 60/056 60/043,670 60/047,501 60/043,578 60/047,614 60/047,593 60/047,589 60/047,585 60/047,588 60/047,599 60/047,595 60/056, 60/056,845 60/056,874 60/056,911 60/056,880 60/056,879 60/056,862 60/056,632

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REFILING DATE: 1997-04-11
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GENERAL INFORMATION:
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Best Local
                                                                                                                         APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-CCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
COUNTAINES: Details PC-DOS/MS-DOS
                                                                      FILING DATE: 06-MAY-PRIOR APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: Human Telomerase Catalytic Subunit
                  PRIOR APPLICATION DATA:
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CITY: San Francisco
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                                    APPLICATION NUMBER: FILING DATE: 09-MAY
                                                                                                              APPLICATION NUMBER:
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ER FILING DATE: 1997-09-05
ER APPLICATION NUMBER: 60/056,884
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/057,669
ER FILING DATE: 1997-09-05
ER APPLICATION NUMBER: 60/049,610
ER FILING DATE: 1997-06-13
ER APPLICATION NUMBER: 60/061,060
ER FILING DATE: 1997-10-02
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Harley, Calvin B.
Andrews, William H.
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Nakamura, Toru
Chapman, Karen B.
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NUMBER:
                                      09-MAY-1997
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100.0%; Pr
                                                      US 08/854,050
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US 08/911,312
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Pred. No.
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US-09-149-476-309/c
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Best Local Similarity 100.
Matches 21; Conservative
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TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 2176 base pairs
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EARLIER APPLICATION NUMBER: 60/040,162
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                    EARLIER
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LOCATION: 1..2176
OTHER INFORMATION:
FEATURE:
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REFERENCE/DOCKET NUMBER: 01:
TELECOMMUNICATION INFORMATION:
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NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,4:
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APPLICATION NUMBER:
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TOPOLOGY: 11
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 APPLICATION NUMBER: 60/040,163
APPLICATION NUMBER: 60/040,163
APPLICATION 1997-03-07
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APPLICATION NUMBER: 60/040,333
FILING DATE: 1997-03-07
                                               APPLICATION NUMBER: 60/040,336 FILING DATE: 1997-03-07
                                                                                    APPLICATION NUMBER: 60/040,334 FILING DATE: 1997-03-07
                                                                                                                     APPLICATION NUMBER: 60/038,621
FILING DATE: 1997-03-07
APPLICATION NUMBER: 60/040,626
FILING DATE: 1997-03-07
                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/US98/04493
FILING DATE: 1998-03-06
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APPLICATION NUMBER: 60/047,600
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14-AUG-1997
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ER FILLING DATE: 1997-04-11
ER FILLING DATE: 1997-04-11
ER FILLING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,674
ER FILLING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,669
ER FILLING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,312
ER FILLING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,313
ER FILLING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,672
ER FILLING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,315
ER FILLING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,315
ER FILLING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/043,315
ER APPLICATION NUMBER: 60/048,974
ER FILLING DATE: 1997-06-06
ER APPLICATION NUMBER: 60/056,886
ER FILLING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,877
ER FILLING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,877
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ER APPLICATION NUMBER: 60/056,877 R APPLICATION NUMBER: 60/047,596
R FILING DATE: 1997-05-23
R APPLICATION NUMBER: 60/047,612
R FILING DATE: 1997-05-23
R APPLICATION NUMBER: 60/047,632
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R FILING DATE: 1997-05-23
R APPLICATION NUMBER: 60/047,601 PR APPLICATION NUMBER: 60/047,617
ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,618
ER FILING DATE: 1997-05-23
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R APPLICATION NUMBER: 60/00
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R FILING DATE: 1997-05-APPLICATION NUMBER: 60/1 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/1 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/043,569 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/ FILING DATE: 1997-04-11 FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/043,580
FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/ FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/ FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1997-05 APPLICATION NUMBER: 60/ FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/043,568 FILING DATE: 1997-04-11 APPLICATION NUMBER: DATE: 1997-05-23 : 1997-05-1997-05-23 1997-05-23 1997-05-60/047,598 60/047,492 60/047,633 60/043,314 60/047,582 60/047,613 60/047,587 60/047,500 60/047,584 60/047,581 60/047,583 60/047,502 60/047,597 60/047,615 EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER
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ER APPLICATION NUMBER: 60/056,631
ER FILLING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,845
ER APPLICATION NUMBER: 60/056,892
ER APPLICATION NUMBER: 60/057,761
ER FILLING DATE: 1997-08-22
ER FILLING DATE: 1997-08-23
ER APPLICATION NUMBER: 60/047,595
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,599
ER APPLICATION NUMBER: 60/047,586
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,585
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,585
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,586
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,590
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,590
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ER APPLICATION NUMBER: 60/047,590
ER FILLING DATE: 1997-05-23 ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,911
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,636
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,874
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,910
ER FILING DATE: 1997-08-22
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ER FILING DATE: 1997-08-22
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ER FILING DATE: 1997-08-22
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ER FILING DATE: 1997-08-26
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ER FILING DATE: 1997-08-26 R FILING DATE: 1997-08-22
IR APPLICATION NUMBER: 60/05
IR FILING DATE: 1997-08-22
IR APPLICATION NUMBER: 60/05
IR FILING DATE: 1997-08-22 ER APPLICATION NUMBER: 60/047,501
ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/043,670
ER FILING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/056,632
ER FILING DATE: 1997-08-22 R APPLICATION NUMBER: 60/04
R FILING DATE: 1997-05-23
R APPLICATION NUMBER: 60/04
R FILING DATE: 1997-05-23
R APPLICATION NUMBER: 60/04
R FILING DATE: 1997-04-11
R FILING DATE: 1997-04-11
R FILING DATE: 1997-04-11 R FILING DATE: 1997-08-22
R FILING DATE: 1997-08-22
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R APPLICATION NUMBER: 60/0:
R FILING DATE: 1997-08-22 R FILING DATE: 1997-08-R APPLICATION NUMBER: 6 R FILING DATE: 1997-08-R APPLICATION NUMBER: 6 FILING DATE: 1997-05-22
APPLICATION NUMBER: 60/056,903
APPLICATION NUMBER: 60/056,808 APPLICATION: NUMBER: 60/056,880 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,894 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60 APPLICATION NUMBER: 6 FILING DATE: 1997-08-NUMBER: 60/043,576: 1997-04-11 NUMBER: 60/047,589: 1997-05-23 1997-08-1997-08-60/043,578 60/047,614 60/047,593 60/056,879 60/056,888 60/056,637 60/056,882 60/056,872 60/056,662 60/056,893

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EARLIER FILING DATE: 1997-08-22
EARLIER APPLICATION NUMBER: 60/056,875
EARLIER FILING DATE: 1997-08-22
EARLIER APPLICATION NUMBER: 60/056,862
EARLIER FILING DATE: 1997-08-22
EARLIER APPLICATION NUMBER: 60/056,867
EARLIER APPLICATION NUMBER: 60/056,887
EARLIER APPLICATION NUMBER: 60/056,887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 46, Application US/08686878A Patent No. 5708157
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EARLIER APPLICATION NUMBER: 60/056,884
EARLIER FILING DATE: 1997-08-22
EARLIER APPLICATION NUMBER: 60/057,669
EARLIER FILING DATE: 1997-09-05
EARLIER APPLICATION NUMBER: 60/049,610
EARLIER FILING DATE: 1997-06-13
EARLIER APPLICATION NUMBER: 60/061,060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EARLIER FILING DATE: 1997-10-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EARLIER
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                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Evans, Cheryl
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES
TITLE OF INVENTION: ENCODING THEM
NUMBER OF SEQUENCES: 71
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   605 TGNAAAAAAAAAAAAAAAA 625
                                                                                 APPLICATION NUMBER: FILING DATE:
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les 21; Conserv
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ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/048,964
ER FILING DATE: 1997-06-06
ER APPLICATION NUMBER: 60/057,650
ER FILING DATE: 1997-09-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/056,876 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,881 FILING DATE: 1997-08-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/056,664
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                                                                                                                                                                                                                                                                                                              87 CambridgePark Drive
                                                                                                                                                                                                                                                                U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LaVallie, Edward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           McCoy, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jacobs, Kenneth
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                                                                                                                                                                                                                                                                                                                                         Genetics Institute, Inc.
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                                                                                                US/08/686,878A
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; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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APPLICANT: MCCOy, John M.
APPLICANT: MCCOy, John M.
APPLICANT: LaVallie, Edward R.
APPLICANT: Collins-Racie, Lisa A.
APPLICANT: Evans, Cheryl
APPLICANT: Evans, Cheryl
APPLICANT: Merberg, David
APPLICANT: Mi, Sha
APPLICANT: Mi, Sha
APPLICANT: Mi, Sha
APPLICANT: Mi, Sha
APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REFERENCE: 6006B,AJ172A
CURRENT APPLICATION NUMBER: US/09/175,928A
CURRENT APPLICATION NUMBER: US/09/175,928A
CURRENT FILING DATE: 1998-10-20
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 45
LENGTH: 265
TYPE: DNA
CDCANTCM: Miss missculus
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US-09-175-928-45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
LENGTH: 265 base pairs
TENGTH: 154 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 45, Approx 8 12921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 3.2%; Score 20; DB 1; Best Local Similarity 100.0%; Pred. No. 4.9; Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: unsure
LOCATION: (67)
                                                NAME/KEY:
LOCATION:
FEATURE:
                          NAME/KEY: unsure
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Mus musculus FEATURE:
FEATURE:
           COCATION: (133)
                                                                                                       FEATURE
                                                                                                                    LOCATION:
                                                                                                                                         NAME/KEY:
                                                                                                                                                           FEATURE:
                                                                                                                                                                                                          FEATURE:
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STRANDEDNESS: double
TOPOLOGY: linear
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                                                               unsure
(121)
                                                                                                                  unsure
(111)
                                                                                                                                                                       unsure
(104)
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NAME/KEY: unsure LOCATION: (175)

EATURE

OCATION: (183)

NAME/KEY: unsure LOCATION: (199)..(200)

NAME/KEY: OCATION: NAME/KEY:

unsure (211)

unsure (223)

unsure (208)

NAME/KEY: unsure LOCATION: (192)

OCATION: AME/KEY: EATURE

(187) unsure

EATURE

PEATURE

NAME/KEY: unsure LOCATION: (172)

OCATION: (164) WAME/KEY: unsure NAME/KEY: unsure LOCATION: (157)

NAME/KEY: unsure

FEATURE: NAME/KEY:

unsure (136)

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APPLICANT: Fisher, Paul B.
TITLE OF INVENTION: DISPLAY
TITLE OF INVENTION: DISPLAY
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham LLP
1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                        Score 20; DB 4;; Pred. No. 4.9; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                          0; Indels
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                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 10
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; Patent No. 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/032,684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 55
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 278-0400
                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 183
CORRESPONDENCE ADDRESS:
ADDRESSEE: TESTA, HURWITZ & THIBEAULT
STREET: High Street Tower - 125 High Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
TITLE OF INVENTION: TO ALZHEIMER'S DISEASE
NUMBER OF SEQUENCES: 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 296 base pairs
                                                                                                                                   APPLICATION NUMBER: US/08/967,101 FILING DATE: 10-NOV-1997 CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
     NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
TELEPAX: (617) 248-7100
                                                                                                                                                                                                                SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                            FILING DATE:
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U
ZIP: 10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    60 GNAAAAAAAAAAAAAAAA 41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                            STATE: Massachusetts
COUNTRY: U.S.A.
ZIP: 02110
                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                     Boston
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100.0%; Pred. No. 4.
tive 0; Mismatches
                                                                                                                      08/592,541
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; NAME/KEY: unsure ; LOCATION: (246) US-09-175-928-45

FEATURE:

NAME/KEY: unsure LOCATION: (243)..(244)

OCATION:

unsure (227)

unsure (225)

Query Match 3.2%; So Best Local Similarity 100.0%; F Matches 20; Conservative 0;

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RESULT 9
US-09-032-684-13/c
; Sequence 13, Application US/09032684
; Patent No. 5882874

GENERAL INFORMATION:

STREET:

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US-09-124-698-82/c
; Sequence 82, Application US/09124698
; Patent No. 6117978
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                                                               RESULT 12
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US-08-592-541-82/c
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                                                                                                                                                                                                                                                                                           TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 82
SEQUENCE CHARACTERISTICS:
LENGTH: 500 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                               Query Match
                                                                                                                                                                                    Matches
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: TESTA, HURWITZ & THIBEAULT
STREET: High Street Tower - 125 High Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                             606 GNAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA (genomic)
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                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                  linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENETIC SEQUENCES AND PROTEINS RELATED
                                                                                                                                                                                           3.2%; Score 20; DB 2;
L00.0%; Pred. No. 4.7;
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                                                                                                                                                                                                                                                                                                                                                                                   Sequence 82, Application US/09127480 Patent No. 6194153 GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                           APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
TITLE OF INVENTION: TO ALZHEIMER'S DISEASE
            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: TESTA, HU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Pitcher, Edmund R. TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 248-7000
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ATTORNEY/AGENT INFORMATION:
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APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, JOHANNA M
APPLICANT: FRASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
NUMBER OF SEQUENCES: 183
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       34 GNAAAAAAAAAAAAAAAA 15
APPLICATION NUMBER:
                                                                                                                                                      COUNTRY:
                                                                                                                                                                        STATE: Massachusetts
                                                                                                                                                                                          STREET: High
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 500 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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High Street Tower - 125 High Street
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High Street Tower - 125 High Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.2%; Score 20; DB 3; Length 500; 100.0%; Pred. No. 4.7;
US/09/127,480
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US-08-496-841C-82/c
Sequence 82, Application US/08496841C
Patent No. 6210919
GENERAL INFORMATION:
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                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 82: US-08-496-841C-82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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 Best Local Similarity
                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (617) 248-7100 INFORMATION FOR SEQ ID NO: 82:
                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 82:
SEQUENCE CHARACTERISTICS:
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NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
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STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ADDRESSEE: Darby & Darby, PC
                                                                                       TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 35,135
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Paul F. Fehlner, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
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                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/496,841C FILING DATE: 28-Jun-1995 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: New York
STATE: New York
                                                                                                                                       TYPE: nucleic acid
                                                                                                                                                                                                                 TELEPHONE: (212) 527-7700
TELEFAX: (212) 753-6237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 805 Third Avenue
                                                                                                                         STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              500 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ROMMENS, JOHANN
FRASER, PAUL E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GEORGE-HYSLOP, PETER H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.2%; UT
100.0%; Pre
3.2%;
100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JOHANNA M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 20;
Pred. No.
Score 20; DB 4; Pred. No. 4.7;
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                   Length 500;
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RESULT 15
US-09-124-523-82/c
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                                                                                                                                                                                                                                                                           В
                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-124-523-82
                                                                                                                                                                                                     US-09-020-956-32/c
                                                                                                                                                                                                                   RESULT 16
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                                                                                                                                                               Sequence 32, Application US/09020956 Patent No. 6261562
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                                                                     GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillin, Davin C.
APPLICANT: Dillin, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS
NUMBER OF SEQUENCES: 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (617) 248-7100 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
TITLE OF INVENTION: TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA: APPLICATION NUMBER:
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                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                             34 GNAAAAAAAAAAAAAAAA 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
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STREET: 6300 (CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                   Match 3.2%; Score 20; DB 4; Length 500; Local Similarity 100.0%; Pred. No. 4.7;
                 ADDRESSEE: SEED and BERRY LLP STREET: 6300 Columbia Center,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   500 base pairs
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                                                                                                                                                                                                                                                                                                                                                  Conservative
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                     701 Fifth Avenue
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US-09-030-607-32/c
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GENERAL INFORMATION:
                         RAME: Mak1, Dav1d J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210121.427C3
TELECOMMUNICATION INFORMATION:
TELEFACAN: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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LENGTH: 789 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: Linear
MOLECULE TYPE: CDNA
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REFERENCE/DOCKET NUMBER: 210
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 32:
                                                                                                                                                                      APPLICATION NUMBER: US/09
FILING DATE: 25-FEB-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Xu, Jiangchun APPLICANT: Xu, Jiangchun APPLICANT: Xu, Jiangchun C. TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FOR NUMBER OF SEQUENCES: 224
                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                   LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 6300 (CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: SEED and BERRY LLP STREET: 6300 Columbia Center, 701 Fifth Avenue
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Local Similarity 100.0%; P
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                                                                                                                                                                                                                                                                                                                                                                                                  98104
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nucleic acid
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            789 base pairs
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                                                                                                                                                                                                                                                       US/09/030,607
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US-09-439-313-32/c
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GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun

APPLICANT: Dillon, Davin C.

APPLICANT: Mitcham, Jennifer L.

APPLICANT: Harlocker, Susan Louise

APPLICANT: Jiang Yuqui

APPLICANT: Reed, Steven G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-605-785-32/c
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                                                                                                                          Sequence 32, Application US/09439313 Patent No. 6329505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Wang, Aijun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 20121. 427C16
CURRENT APPLICATION NUMBER: US/09/605,785
CURRENT FILING DATE: 2000-06-27
NUMBER OF SEQ ID NOS: 835
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                          Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                 606 GNAAAAAAAAAAAAAAAA 625
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TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                              ch 3.2%; Score 20; DB 4; Length 789; I Similarity 100.0%; Pred. No. 4.5; 20; Conservative 0; Mismatches 0; Indels
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6321716
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Carter, Darrick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vedvick,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Li, Samuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kalos, Michael D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Henderson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jiang, Yuqu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Harlocker, Susan L.
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ick, Thomas S.
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100.0%; Pred. No. 4.5;
Pative 0; Mismatches
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                          US-09-232-149A-32/c; Sequence 32, Application US/09232149A; Patent No. 6465611;
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                                                                                            RESULT 21
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: COMPOSITIONS OF PROSTATE CANCER
TITLE OF INVENTION: DARANOSIS OF PROSTATE CANCER
TITLE OF INVENTION: COMPOSITION OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT FILING DATE: 1999-11-12
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Dillon, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.427C8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 32, Application US/09352616A Patent No. 6395278
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                GENERAL
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APPLICANT: Xu, Jiangchun
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LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
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ORGANISM: Homo sapien
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TYPE: DNA
ORGANISM: Homo sapien
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NAME/KEY: misc_feature
LOCATION: (1)...(789)
OTHER INFORMATION: n =
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                  INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FastSEQ for Windows Version 3.0
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Fanger, Gary
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100.0%; Pred. No.
tive 0; Mismatch
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100.0%; Pred. No. 4.1
ive 0; Mismatches
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APPLICANT: Dillon, Davin C.

APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE
TITLE OF INVENTION: CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.427C6
CURRENT APPLICATION NUMBER: US/09/232,149A
CURRENT FILING DATE: 1999-01-15
NUMBER OF SEQ ID NOS: 338
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/149,476
CURRENT FILING DATE: 1998-09-08
EARLIER APPLICATION NUMBER: PCT/US98/04493
EARLIER FILING DATE: 1998-03-06
EARLIER APPLICATION NUMBER: 60/040,162
EARLIER FILING DATE: 1997-03-07
EARLIER FILING DATE: 1997-03-07
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                                                                                                                                                                                                                                                     EARLIER APPLICATION NUMBER: 60/047,615
EARLIER FILING DATE: 1997-05-23
EARLIER APPLICATION NUMBER: 60/047,597
EARLIER FILING DATE: 1997-05-23
EARLIER APPLICATION NUMBER: 60/047,502
EARLIER FILING DATE: 1997-05-23
                                                                                                                                                                                                                                                                                                                                                                                               EARLIER APPLICATION NUMBER: 60/040,336
EARLIER FILING DATE: 1997-03-07
EARLIER APPLICATION NUMBER: 60/040,163
EARLIER FILING DATE: 1997-03-07
EARLIER APPLICATION NUMBER: 60/047,600
EARLIER FILING DATE: 1997-05-23
                                                             EARLIER APPLICATION NUMBER: 60/047,617
EARLIER FILING DATE: 1997-05-23
EARLIER APPLICATION NUMBER: 60/047,618
EARLIER FILING DATE: 1997-05-23
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EARLIER FILING DATE: 1997-03-07
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LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
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ORGANISM: Homo sapien
FEATURE:
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                                                                                                                                                            FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,583
FILING DATE: 1997-05-23
              APPLICATION NUMBER: 60/047,503 FILING DATE: 1997-05-23
                                                                                                                                                                                                                                 APPLICATION NUMBER: 60/047,633
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APPLICATION NUMBER: 60/047,592
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ER FILING DATE: 1997-08-22
ER FILING DATE: 1997-08-22
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ER APPLICATION NUMBER: 607056,882
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 607056,637
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 607056,903
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 607056,903
ER FILING DATE: 1997-08-22 PR FILING DATE: 1997-04-11

PR APPLICATION NUMBER: 60/043,315

PR FILING DATE: 1997-04-11

PR APPLICATION NUMBER: 60/048,974

PR FILING DATE: 1997-06-06

PR APPLICATION NUMBER: 60/056,886

PR APPLICATION NUMBER: 60/056,886

PR FILING DATE: 1997-08-22 ER APPLICATION NUMBER: 60,047,632
ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60,047,601
ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60,043,580
ER FILING DATE: 1997-04-11
ER APPLICATION NUMBER: 60,043,568
ER FILING DATE: 1997-04-11
ER APPLICATION NUMBER: 60,043,568
ER FILING DATE: 1997-04-11 FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,893
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,630
FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/L APPLICATION NUMBER: 60/056,878 FILING DATE: 1997-08-22 FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/043,312
FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/043,313
FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/ FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/043,672 FILING DATE: 1997-04-11 FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/043,674
FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,669 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,314
FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/043,569
FILING DATE: 1997-04-11 APPLICATION NUMBER: APPLICATION NUMBER: 60/043,311 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/ FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,598 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,613 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/ FILING DATE: 1997-05-23 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/047,612 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,596 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/ FILING DATE: 1997-05-23 APPLICATION NUMBER: FILING DATE: 1997-0 APPLICATION NUMBER: 60 FILING DATE: 1997-05-FILING DATE: 1997-05-APPLICATION NUMBER: 61FILING DATE: 1997-05-1997-05-23 60/056,889 60/056,877 60/043,671 60/047,582 60/047,500 60/047,492 60/047,587 60/047,584 60/047,581

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R APPLICATION NUMBER: 60/01
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R FILING DATE: 1997-05-23
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R APPLICATION NUMBER: 60/043,576
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ER FILLING DATE: 1997-05-23
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ER FILLING DATE: 1997-05-23
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ER APPLICATION NUMBER: 60/047,593
ER APPLICATION NUMBER: 60/047,593 APPLICATION NUMBER: 60/056,887 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,908 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/048,964 APPLICATE: 1997-00-22
APPLICATION NUMBER: 60/C
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APPLICATION NUMBER: 60/043,670
TITING DATE: 1997-04-11 FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/056,632
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,664
FILING DATE: 1997-08-22 FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,588
FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,501 FILING DATE: 1997-05-23 APPLICATE: 1997-08-22
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,864
APPLICATION NUMBER: 60/056,663 APPLICATION NUMBER: 60/ FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/ FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/ FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,888 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,879 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/057,761 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,892 FILING DATE: 1997-08-22 FILING DATE: 1997-0
APPLICATION NUMBER: APPLICATION NUMBER: 60/0 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,874 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,636 APPLICATION NUMBER: 60/0 FILING DATE: 1997-08-22 APPLICATION NUMBER: APPLICATION NUMBER: 60/056,880 FILING DATE: 1997-08-22 FILING DATE: 1997-08-60/056,862 60/056,875 60/056,909 60/047,599 60/047,595 60/056,845 60/056,631 60/056,911 60/056,894

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Best Local :
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LENGTH: 1210
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EARLIER APPLICATION NUMBER: 60/057,650
EARLIER FILING DATE: 1997-09-05
EARLIER APPLICATION NUMBER: 60/056,884
EARLIER FILING DATE: 1997-08-22
EARLIER APPLICATION NUMBER: 60/057,669
EARLIER FILING DATE: 1997-09-05
EARLIER FILING DATE: 1997-09-05
                                                                                                                                                                                                                                                 Sequence 4, Application US/09648040 Patent No. 6436665
                                                                                                                                                                                                                             GENERAL INFORMATION:
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SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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CURRENT FILING DATE: 1999-11-18
PRIOR APPLICATION NUMBER: 60/109,283
PRIOR ETLING DATE: 1998-11-20
NUMBER OF SEQ ID NOS: 33
SOFTWARE: Microsoft Office 97
                                 APPLICANT: Robert G. Kulmelis
TITLE OF INVENTION: METHODS FOR CODING AND SORTING IN VITRO
TITLE OF INVENTION: TRANSLATED PROTEINS
FILE REFERENCE: 50036/032002
CURRENT APPLICATION NUMBER: US/09/648,040
CURRENT FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/151,261
PRIOR FILING DATE: 1999-08-27
NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Famodu, Omolayo O. APPLICANT: Orozco, Buddy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EARLIER FILING DATE: 1997-10-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Sterol Metabolism Enzymes
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                      SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: unsure
LOCATION: 1129
OTHER INFORMATION: any nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 h 3.2%; Score 20; DB 4; Length 1210; Similarity 100.0%; Pred. No. 4.4;
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Rafalski, Antoni
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; NAME/KEY: misc_feature
; LOCATION: 40
; OTHER INFORMATION: /product= "N is A, C or G"
US-08-859-106A-4
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                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 196:
FILING DATE: 22-MAY-1996
ATTORNEY/AGENT INFORMATION:
NAME: MCGOWAN, MAlcolm K.
REGISTRATION NUMBER: 39,300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature LOCATION: 10 OTHER INFORMATION: n at position 10 can be a, t, c, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Encoding molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
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                                                                                                                                                                                                   TELEFAX: (703) 836-2021 INFORMATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
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APPLICANT:
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TITLE OF INVENTION: ASPERGILLUS BY RECOMBINANT METHODS
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                                                                     FEATURE:
                                                                                   ANTI-SENSE:
                                                                                                                    MOLECULE TYPE:
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                                                                                                   HYPOTHETICAL:
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                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE:
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                                                                                                                                                                 nucleotide
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                                                                                                                                                                                     40 base pairs
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NGUYEN, Quoc Khanh
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Conservative 0;
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                                                                                                   genomic DNA
No
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20-MAY-1997
                                                                                                                                                    single-stranded
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100.0%; Pred. No.
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CURRENT APPLICATION NUMBER: US/09/011,540
CURRENT FILING DATE: 1998-04-20
EARLIER APPLICATION NUMBER: DE 19526485.1
EARLIER FILING DATE: 1995-07-20
EARLIER APPLICATION NUMBER: PCT/EP96/01430
EARLIER FILING DATE: 1996-04-01
NUMBER OF SEQ ID NOS: 12
SOFTMARE: Patentin Ver. 2.0
SEQ ID NO 11
SEQ ID NO 11
LENGTH: 40
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          SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 5
LENGTH: 51
                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                   APPLICANT: WEINDEL, KUTT
APPLICANT: SEIDEL, Cristoph
APPLICANT: LASSONCZYK, Gerhard
TITLE OF INVENTION: METHOD OF DETECTING A SUBSTANCE TO BE ANALYZED
FILE REFERENCE: 101614-07096
CURRENT APPLICATION NUMBER: US/09/051,079A
CURRENT FILING DATE: 1998-09-10
EARLIER APPLICATION NUMBER: PCT/EP96/04358
EARLIER APPLICATION NUMBER: DE/195 37 952.7
EARLIER FILING DATE: 1996-10-12
EARLIER FILING DATE: 1995-10-12
EARLIER FILING DATE: 1995-10-12
EARLIER FILING DATE: 1995-10-12
EARLIER FILING DATE: 1995-10-12
EARLIER FILING DATE: 1995-10-12
EARLIER FILING DATE: 1995-10-12
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APPLICANT: GOTTSCHALK, Michael
APPLICANT: KHANH, Nguyen Quoc
APPLICANT: WOLF, Sabine
APPLICANT: WOLF, Sabine
APPLICANT: PLAINER, Hermann
TITLE OF INVENTION: LEUCINE AMINOPEPTIDASES PRODUCED RECOMBINANTLY FROM
TITLE OF INVENTION: ASPERGILLUS SOYAE
FILE REFERENCE: 015200-051
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PEATURE:
NAME/KEY: misc_feature
LOCATION: (39)...(40)
OTHER INFORMATION: N is A, C or G.
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100.0%; Pred. No.
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Pred. No.
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US-08-618-100B-9/c
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                      REFERENCE/DOCKET NUMBER: 32,327
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                      OVERANTING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/618,100B
FILING DATE: March 19, 1996
CLASSIFTOATTOM. F1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                             FILING DATE: October 30, 1995
APPLICATION NUMBER: 08/510,584
FILING DATE: August 2, 1995
APPLICATION NUMBER: 08/418,096
FILING DATE: APPLI 5, 1995
APPLICATION NUMBER: 08/408,584
FILING DATE: March 20, 1995
                                                                                                                                                                                                                                                                            CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial OTHER INFORMATION: Oligodesoxyribonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: SULCCITY: LOS Angeles CALIFORNIA "S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Croston, Glenn L.
APPLICANT: Miller, Stephen G.
APPLICANT: MILLER, Stephen G.
APPLICANT: MODULATORS OF Ob GENE AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: n means DNP bound via DNP-TEG
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                                                                                                                                                                                                               APPLICATION NUMBER:
FILING DATE: October
APPLICATION NUMBER:
FILING DATE: August
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                607 NAAAAAAAAAAAAAAAAA 625
                   TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                    NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: 3.5" Di
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION:
                                                                                                             Warburg, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/08618100B
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633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Auwerx, Johan
de Vos, Piet
Staels, Bart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Briggs, Michael R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                               IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Description of Artificial Sequence:
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                                                                                                                                                                                                                                                                08/558,588
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RESULT 29 US-08-702-344-7

Sequence 7, Appl patent No. 57233

Application US/08702344

GENERAL INFORMATION:

APPLICANT:

APPLICANT:
APPLICANT:

McCoy, John Lavallie, Edward

John

Jacobs, Kenneth

Racie, Lisa

NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:

APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES
TITLE OF INVENTION: ENCODING THEM

US-08-618-100B-9

FEATURE:

OTHER INFORMATION: "N" represents any base

TOPOLOGY: linear

SEQUENCE CHARACTERISTICS: LENGTH: 52 base pairs TYPE: nucleic acid STRANDEDNESS: single

Matches Query Match

Local Similarity

3.0%;

Score 19; pred. No.

DB 3; Length 52; 15;

Conservative

0;

Mismatches

0; Indels

0;

607 NAAAAAAAAAAAAAAAA 625

51 NAAAAAAAAAAAAAAA 33

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Gaps
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US-08-702-344-22
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US-09-269-911A-7/c
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                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                Sequence 7, Application US/09269911A Patent No. 6228589
                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                         SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                       Best
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INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                                                                                                                                   APPLICANT: Brenner, Sydney
TITLE OF INVENTION: MEASUREMENT OF GENE EXPRESSION PROFILES
TITLE OF INVENTION: IN TOXICITY-DETERMINATION
FILE REFERENCE: 5525-0025.10
                                                                                                       CURRENT APPLICATION NUMBER: US/09/269,911A CURRENT FILING DATE: 2000-02-28
                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PRI-DASE #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: SECRETED PROTITIVE OF INVENTION: ENCODING THEM
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           TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
FEATURE:
                                          ENGTH: 69
                                                                                                                                                                                                                                                                                                                                        607 NAAAAAAAAAAAAAAAAA 625
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE:
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                                                                                                                                                                                                                                                                                                            22 NAAAAAAAAAAAAAAAA 40
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                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleic acid
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                                                                           FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Merberg, David
Merberg, Maurice
Treacy, Maurice
Spaulding, Vikki
Spaulding, Vikki
Spaulding, Portice
Spaulding, Them
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    69 base pairs
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                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (617) 498-8224
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     double
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100.0%; Pr
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Pred. No.
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В Ş US-08-702-344-7 Query Match MOLECULE TYPE: cDNA 607 NAAAAAAAAAAAAAAAAA 625 Local Similarity es 19; Conserv 5 NAAAAAAAAAAAAAAAA 23 Conservative 3.0%; but 100.0%; Pr/ ''' 0; Score 19; pred. No. Mismatches DB 1; Length 69; 15;

0

TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
TELEPHONE: (617) 876-5851
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 69 base pairs

TYPE: nucleic acid STRANDEDNESS: double

TOPOLOGY: 11

linear

ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724

CURRENT APPLICATION DATA:

APPLICATION NUMBER:

FILING DATE: CLASSIFICATION: 536

COMPUTER: IBM PC COMPATIBLE
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Pelana

PatentIn Release #1.0, Version #1.30

US/08/702,344

0;

COMPUTER READABLE FORM:

02140

COUNTRY:

U.S.A.

STREET: ADDRESSEE:

87 CambridgePark Drive

Genetics Institute, Inc.

Cambridge

Massachusetts

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                                                                                                                        US-08-664-596B-3
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                                                              Matches
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Best Local :
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INFORMATION FOR SEQ ID NO: 3:
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NAME/REY: misc_feature
LOCATION: (65)...(65)
OTHER INFORMATION: n = cDNA from library
                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/664,596B
FILING DATE:
CLASSIFICATION: 514
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APPLICANT:
                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                            NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Bowman, Michael
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES
TITLE OF INVENTION: ENCODING THEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                            607 NAAAAAAAAAAAAAAAAA 625
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                                                                           Local
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8 NAAAAAAAAAAAAAAAA 26
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5. 5807703
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                                                                         Similarity
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                                                                                                                                                                                nucleic acid
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                                                                                                                                                                                               84 base pairs
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Treacy, Maurice
Evans, Cheryl
Spaulding, Vikki
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LaVallie, Edward
                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genetics Institute, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                                                                   double
                                                                         3.0%; Score 19;
100.0%; Pred. No.
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                                                         Mismatches
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                                                                         DB 1;
15;
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                                                                                     Length 84;
                                                         Indels
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; TOPOLOGY: li
; MOLECULE TYPE:
US-08-738-367-3
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US-08-991-789A-266/c
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US-08-738-367-3
                                                                                       Sequence 266, Application US/08991789A
Patent No. 6225054
GENERAL INFORMATION:
SMITH, John M.
SMITH, John M.
Reed, Steven G.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TREATMENT AND DIAGNOSIS OF BREAST CANCER
                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (617) 876-5851 INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/738,367
FILING DATE:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 84 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION: NAME: Brown, Scott A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES
TITLE OF INVENTION: ENCODING THEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                  607 НААААААААААААААААА 625
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                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 87 Cambridge STATE: Massachus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Brown, Scott A. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                   8 NAAAAAAAAAAAAAAAA 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                             NUMBER OF SEQUENCES: 292
CORRESPONDENCE ADDRESS EE Seed IP Law Group
STREET: 701 Fifth Avenue, Suite 6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid
 CITY: Seattle
STATE: Washing
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                                                                                                                                                                                                                                                                                                                                                                     Conservative
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LaVallie, E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            U.S.A.
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Washington
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                                                                                                                                                                                                                                                                                                                                                                                   3.0%; Score 19;
100.0%; Pred. No.
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COUNTRY: USA

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US-09-062-451-266/c
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                                         TELEFAX: (206) 682-60 INFORMATION FOR SEQ ID NO:
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INFORMATION FOR SEQ ID NO: 266:
SEQUENCE CHARACTERISTICS:
LENGTH: 193 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                              NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION MABER: US/19/062,451
FILING DATE: 04-APR-1997
                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 297
CORRESPONDENCE ADDRESS:
ADDRESSE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Frudakis, Tony N. APPLICANT: Smith, John M. APPLICANT: Reed, Steven G.
                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                     CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF BREAST CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              180 NAAAAAAAAAAAAAAAAA 162
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Local Similarity 100.0%; F
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PSteptin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 210121.419C3 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/991,789A
FILING DATE: 11-Dec-1997
CLASSIFICATION: <UNKnown>
ATTORNEY/AGENT INFORMATION:
NAME: POTTER, Jane E. R.
REGISTRATION NUMBER: 33,332
REGISTRATION NUMBER: 33,332
                                                                                                                                                                                                                                                                                                                                                                                                        98104-7092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seattle
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193 base pairs
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266:
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; Pred. No.
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                                                                                                                            US-08-686-878A-37
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Patent No. 5708157
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                                                           Query Match
Best Local Similarity
Matches 19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Evans, Cheryl
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEI
TITLE OF INVENTION: ENCODING THEM
NUMBER OF SEQUENCES: 71
                                                                                                                                                                                                                                     TELEFAX: (617) 876-58
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc
STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                 NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                         MOLECULE TYPE:
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LENGTH: 208 base pairs
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                             607 NAAAAAAAAAAAAAAAA 625
161 NAAAAAAAAAAAAAAAAA 179
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STRANDEDNESS: sing
TOPOLOGY: linear
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Local Similarity 100.0%;
nes 19; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 87 Cambridge
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DEDNESS: double
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(617) 876-5851
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LaVallie, Edward
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                                                             Conservative
                                                                                                                                                            linear
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                                                                            100.0%;
                                                                              3.0%; Score 19; DB
100.0%; Pred. No. 14;
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                                                             0;
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                                                               Mismatches
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. 14;
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                                                                                              Length 208;
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RESULT 37 US-08-967-101-98/c ; Sequence 98, Application US/08967101

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US-08-592-541-98/c
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Best Local Similarity
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GENERAL INFORMATION:
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TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
TITLE OF INVENTION: TO ALZHELMER'S DISEASE
NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 208 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC computable
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                    APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                              TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED TITLE OF INVENTION: TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 159 NAAAAAAAAAAAAAAAA 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   607 NAAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: High Street.7
CITY: Boston
STATE: Massachusetts
                                                                                                                                                   STREET: High
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/01 FILING DATE: 10-NOV-1997
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                                                                                                             COUNTRY:
                                                                                                                                                                                       ADDRESSEE:
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                                                                                                                                   Massachusetts
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                                                                                                                                                                     E: TESTA, HURWITZ & THIBEAULT
High Street Tower - 125 High Street
                                                                                                               U.S.A.
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PatentIn Release #1.0, Version #1.30
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100.0%; Pred. No.
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US-09-124-698-98/c
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                                                   US-09-124-698-98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 98:
SEQUENCE CHARACTERISTICS:
LENGTH: 208 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 98, Application US/09124698 Patent No. 6117978
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               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:

APPLICANT: ST. GEORGE-HYSLOP,

APPLICANT: ROMMENS, JOHANNA M

APPLICANT: FRASER, PAUL E
                                                                                                                                                                   TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 98:
                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: PITCHET, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOTTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/592,541
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: GE
TITLE OF INVENTION: TO
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA (genomic)
                                                               MOLECULE TYPE: DNA (genomic)
                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
TOPOLOGY: linear
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Boston
STATE: Massachusetts
                                                                                     TOPOLOGY: 1i
                                                                                                                   TYPE:
                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/124,698
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                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02110
                                                                                                               nucleic acid
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High Street Tower - 125 High Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U.S.A.
                                                                                   linear
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                                                                                                   single
3.0%;
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Score 19;
Pred. No.
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DB
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                 Length 208;
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В

Matches

19;

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RESULT 41
US-08-496-841C-98/C
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                                                                                                                                             Sequence 98, Application US/08496841C Patent No. 6210919
GENERAL INFORMATION:
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
TITLE OF INVENTION: TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (617) 248-7100 INFORMATION FOR SEQ ID NO: 98:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Pitcher, Edmund R. TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 208 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                              159 NAAAAAAAAAAAAAAAAA 141
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STATE: Massachusette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                 3.0%; Score 19; Local Similarity 100.0%; Pred. No. es 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                     TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED TO ALZHEIMER'S DISEASE
                                                                                                          APPLICANT: ST. GEORGE-HYSLOP, PETER H
                   CORRESPONDENCE ADDRESS
                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02110
ADDRESSEE: Darby & Darby, PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/09127480
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                                                                                             FRASER, PAUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA (genomic)
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125 High Street
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                                                                                                                                                                                                                                                                                                                                                                                         DB 4;
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US-09-175-928-37, Application US/09175928A
: Patent No. 6312921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear;
; MOLECULE TYPE: DNA (genomic)
; SEQUENCE DESCRIPTION: SEQ ID NO: 98:
US-08-496-841C-98
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                                                                                                                                                                                                  SEQ ID NO 37
LENGTH: 208
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                    APPLICANT: Mi, Sha
APPLICANT: Genetics Institute, Inc.
APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REFERENCE: 6006B.AJ172A
FILE REFERENCE: 6006B.AJ172A
                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Evans, Cheryl
APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Mi, Sha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Jacobs, Kenneth APPLICANT: McCoy, John M. APPLICANT: LAVAILIE, Edward R.
                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/175,928A CURRENT FILING DATE: 1998-10-20
                                                                                                                                                                                                                                                         SOFTWARE:
                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
               NAME/KEY:
LOCATION:
                                                                                                                           NAME/KEY:
LOCATION:
                                                                                                                                                                ORGANISM: Mus musculus FEATURE:
                                                   FEATURE:
                                                                                       FEATURE:
NAME/KEY: unsure
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FEATURE:
                                                                      COCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Paul F. Fehlner, Ph.D. REGISTRATION NUMBER: 35,135
TELECOMMUICATION INFORMATION: TELEPHONE: (212) 527-7700
TELEFAX: (212) 753-6237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/496,841C
FILING DATE: 28-Jun-1995
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                       208
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                                                                                                                                                                                                                                                         PatentIn Ver. 2.0
               unsure
(31)
                                                                                                                         unsure
(8)
                                                                    (29)
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OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A. ZIP: 10022
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 805 Third Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 208 base pairs
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14;
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Matches
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LOCATION: (161)
                                                                                                                                                                                                                                                                                                       NAME/KEY: unsure LOCATION: (137)
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LOCATION: (133)
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LOCATION: (130)..(131)
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LOCATION: (100)..(101)
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LOCATION: (91)
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OCATION: (121)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: unsure
                                                                                                                                                                             161 NAAAAAAAAAAAAAAAAA 179
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 STREET:
CITY: E
                                                                                                                                                                                                                    Local Similarity les 19; Conserv
                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                  unsure
(128)
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(119)
Boston
        E: TESTA, HURWITZ & THIBEAULT
High Street Tower - 125 High Street
                                                                                                                                                                                                                       Conservative
                                        GENETIC SEQUENCES AND PROTEINS RELATED TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                    3.0%; Score 19; DB 4; Length 208; 100.0%; Pred. No. 14; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 98:
SEQUENCE CHARACTERISTICS:
LENGTH: 208 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 43 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/686,878A
FILING DATE:
CIASTITICATION. 626
                                                                                                                                                                                                                                                                                                            APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Evans, Cheryl
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES
TITLE OF INVENTION: ENCODING THEM
NUMBER OF SEQUENCES: 71
CORRESSPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                  CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
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STRANDEDNESS: sing
TOPOLOGY: linear
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NAME: Brown, Scott A. REGISTRATION NUMBER: 32,724
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LaVallie, Edward
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APPLICANT: Endege, Wilson O.
APPLICANT: Steinmann, Kathleen E.
APPLICANT: Astle, Jon H.
APPLICANT: Astle, Jon H.
APPLICANT: Burgess, Christopher C.
APPLICANT: Bushnell, Steven E.
APPLICANT: Bushnell, Steven E.
APPLICANT: Carroll III, Eddie
APPLICANT: Carroll III, Eddie
APPLICANT: Carroll III, Eddie
APPLICANT: Carroll Tonam M.
APPLICANT: Ford, Donna M.
APPLICANT: Monahan, John E.
APPLICANT: Schlegel, Robert
TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TITLE OF INVENTION: PRODUCTS
FILE REFERENCE: CCD-257 (US)
CURRENT APPLICATION NUMBER: US/09/328,111
CURRENT FILING DATE: 1999-06-08
EARLIER FILING DATE: 1999-06-08
EARLIER FILING DATE: 1998-06-10
NUMBER OF SEQ ID NOS: 850
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 715
LENGTH: 310
ORGANISM: Homo sapiens
FEATURE:
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US-09-328-111-715
; Sequence 715, Application US/09328111
; Patent No. 6262333
; GENERAL INFORMATION:
                                                                                                                                                                                                                                      FEATURE:

NAME/KEY: misc_feature

LOCATION: (1)...(310)

OTHER INFORMATION: n = A,T,C or G
US-09-328-111-715
Search completed: November 7, Job time: 31.4202 secs
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Matches 19
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TELEPHONE: (617) 498-8224
TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 43:
SEQUENCE CHARACTERISTICS:
LENGTH: 244 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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Maximum DB seq length: 2000000000
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/cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
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) US-09-825-294-214

) US-09-825-294-211

) US-09-825-294-208

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0 US-09-867-701-3375

10 US-09-938-842A-1283

10 US-09-935-299-6936

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RESULT 1 US-09-825-294 / Sequence 21 / Sequence 21 / Patent No. / GENERAL INF / APPLICANT: / APPLICANT: / APPLICANT: / APPLICANT: / APPLICANT:		4	4	c 43	_		4	ωι	w e	w		ω.	w	w	w	31	30	c 29	N	27	c 26	25	N	N	c 22	N	20	
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: LOCATION: (1)...(625)
: OTHER INFORMATION: n = A,T,C or
US-09-825-294-210
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TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 210
LENGTH: 625
TYPE: DIA
ORGANISM: Homo sapiens
FEATURE:
FASTURE: 10 FEATURE:
FEATURE: 10 FEATURE:
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Matches 625; Conserv
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APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOSTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10876
LENGTH: 625
CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC
                                                                    GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS:
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 214
LENGTH: 1897
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(1897)
OTHER INFORMATION: n = A,T,C or G
US-09-825-294-214
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Best Local s
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                           GCCCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                     GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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ilarity 100.0%;
Conservative (
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RESULT 2 US-09-867-701-10876

Sequence 10876, Application Patent No. US20020132237A1

US/09867701

GENERAL INFORMATION:

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; NAME/KEY: misc\_feature ; LCCATION: 607 ; OTHER INFORMATION: n = A,T,C US-09-867-701-10876

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FEATURE:

ORGANISM: Homo sapiens

TYPE: DNA

Query Match Best Local

Matches

al Similarity 625; Conserv

Conservative

Mismatches

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Sequence 205 Application US/09825294

Patent No. US20020004491A1

GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.

APPLICANT: Fling, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA
FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF 520 ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 205

LENCTH: 1619

TYPE: DNA

ORGANISM: HOTO SERVICE
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Best Local Similarity
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99.8%;
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Pred. No. 9.7e-187;
0; Mismatches 0;
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APPLICANT: XU, Jiangchun
APPLICANT: XU, Jiangchun
APPLICANT: XTOlk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Filng, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CF
FILE REFERENCE: 21021 484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 211
LENGTH: 1619
TYPE: DNA
ORGANISM: Homo sapiens
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Best Local S
Matches 606
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                                                                             1052 GCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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CCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGA 239
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Pred. No. 9.7e-187;
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
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 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
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                                                                                                                                                                                                                                                                                                                       Length 1362;
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APPLICANT: Harlocker, Susan L.

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEQ ID NOS: 10912

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 1516

LENGTH: 373

TYPE: DNA

ORGANISM: Homo sapien

US-09-867-701-1516
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                                                                                                                                           Sequence 2409, Application US/09867701 Patent No. US20020132237A1 GENERAL INFORMATION:
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Best Local Similarity
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
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GACAGTG 1598 GACAGTG 606

US-09-825-294-208/c

GENERAL INFORMATION:

APPLICANT: Stolk, John A. APPLICANT: Algate, Paul A. APPLICANT: Fling, Steven P. APPLICANT: Xu, Jiangchun Sequence 208, Application US/09825294 Patent No. US20020004491A1

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Вb 8

1215

; TYPE: DNA ; ORGANISM: Homo sapiens US-09-825-294-208

SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 208 LENGTH: 1362

Query Match Best Local S Matches 359

Local Similarity

Conservative

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1155

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GENERAL INFORMATION:

APPLICANT: Aglate, Paul A.

APPLICANT: Jones, Robert.

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEQ ID NOS: 10912

SOFTWARE: FBStSEQ for Windows Version 4.0

SEQ ID NO 4240

LENGTH: 409
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; ORGANISM: Homo sapien
US-09-867-701-4240
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2409
LENGTH: 349
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Best Local
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315
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                                                                                                                                                                                             Local Similarity
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                                                           ATCAGCCCTGGGCAGACACGAGCAGGAGGGGAGAGACAGAGAAAAAGAAAAACACAGCATGA
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; Pred. No. 9.4e-115;
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                                                                                                                                                                                          Score 263; DB 10; Length 409; Pred. No. 4.5e-97;
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APPLICANT: Aglate, Paul A.

APPLICANT: Jones, Robert

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEQ ID NOS: 10912

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 1532

LENGTH: 390
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; ORGANISM: Homo sapien
US-09-867-701-8894
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APPLICANT: Harlocker, Robert

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION UMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8894
LENGTH: 201
                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                    Sequence 1532, Application US/09867701 Patent No. US20020132237A1
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Best Local
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                    ORGANISM: Homo sapien FEATURE:
NAME/KEY: misc_feature
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; OTHER INFORMATION: n = A,T,C or
US-09-867-701-1532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; NAME/KEY: misc_feature
; LOCATION: (1)...(558)
; OTHER INFORMATION: n = A,T,C
US-09-867-701-2375
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CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2375
LENGTH: 558
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APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapien
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                                                                                                                                                                                                                                                                                                                                                                                                        198 TGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCC 257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              134 ACAGCACCTGCACGAACACCCCGCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGT 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          138 ACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACCACCGTGTACAGGAGCGGGT 197
                                                                                                                                                                                                                                                                                                                                                                                  194 TGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGAGCATCATGTACGCCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CAATTTTTCAGTGTTGGACTTGACAGCTTCTTTT 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CAATTTTTCAGTGTTGGACTTGACAGCTTCTTTT 519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTAC 266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTAC 485
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGACACGAGCAGGAGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAGTAAAT 206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         120;
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 Mismatches

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US-09-783-590-6936
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SEQ ID NO 7826
LENGTH: 381
                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Dillon, Patrick J.
APPLICANT: Haseltine, Willia
APPLICANT: Li, Haodong
APPLICANT: Li, Haodong
APPLICANT: Rosen, Craig A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 5379
SEQ ID NO 1283
LENGTH: 819
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                                                                                                                                                                                                                                                                  Sequence 6936, Application US/09783590 Patent No. US20020110850A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
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APPLICANT: Zhu, Tong
TITLE OF INVENTION: STRESS-REGULATED GENES OF PLANTS, TRANSGENIC PLANTS CONTAINING
TITLE OF INVENTION: SAME, AND METHODS OF USE
FILE REFERENCE: SCRIP1300-3
CURRENT APPLICATION NUMBER: US/09/938,842A
CURRENT FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: US 60/227,866
PRIOR FILING DATE: 2000-08-24
PRIOR FILING DATE: 2000-01-16
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-01-16
                CURRENT APPLICATION NUMBER: US/09/783,590 CURRENT FILING DATE: 2000-02-15 PRIOR APPLICATION NUMBER: 08/420,856 PRIOR FILING DATE: 1995-04-12
                                                                                               APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences,
FILE REFERENCE: PO-16.2C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Harper, Jeff APPLICANT: Kreps, Joel
PRIOR APPLICATION NUMBER: 08/346,731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         715 GCAGGAGGGAGAGACAGAGAAA 736
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nes 22; Conserv
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                                                                                                                                                                                                     Haseltine, William
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100.0%; Pred. No. 2.4e-28
vative 0; Mismatches (
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100.0%; Pred. No.
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Sequence 698, Application US/09925299

Patent No. US20020055627A1

GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA102

CURRENT APPLICATION NUMBER: US/09/925,299

CURRENT FILING DATE: 2001-08-10

PRIOR APPLICATION NUMBER: PCT/US00/05883

PRIOR APPLICATION NUMBER: 60/124,270

PRIOR APPLICATION UNMBER: 60/124,270

PRIOR FILING DATE: 1999-03-12

NUMBER OF SEQ ID NOS: 1556
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Best Local :
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NUMBER OF SEQ ID NOS: 12485
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 6936
LENGTH: 322
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LOCATION: (292)
OTHER INFORMATION: n equals a,t,g,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc feature
LOCATION: (241)
OTHER INFORMATION: n equals a,t,g,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc feature
LOCATION: (236)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (216)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (258)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (283)
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LOCATION: (20)
OTHER INFORMATION: n equals a,t,g,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                   605 TGNAAAAAAAAAAAAAAAA 625
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                                                                                                                                    ; NAME/KEY: misc_feature
; LCCATION: (1)...(547)
; OTHER INFORMATION: n = A,T,C or G
US-09-834-975-379
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                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 379 LENGTH: 547
                                                                   Matches
                                                                            Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 379, Application US/09834975 Patent No. US20020110815A1
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Best Local
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                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/834,975
CURRENT FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 60/197,538
PRIOR FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 698
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Brown, Jeffrey
APPLICANT: Bolt, Andrew
APPLICANT: Wan Huffel, Christophe
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY
TITLE OF INVENTION: OF HUMAN CANCERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lillie, James APPLICANT: Brown, Jeffre
                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: MRI-016B
                                                                                                                                                                                                                  ORGANISM: Homo sapiens
                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
LOCATION: (496)
OTHER INFORMATION: n equals a,t,g,
NAME/KEY: misc_feature
LOCATION: (499)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (480)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (449)
OTHER INFORMATION: n equals a,t,g,
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ORGANISM: Homo
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LOCATION: (358)
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527 TGNAAAAAAAAAAAAAAAA 547
                605 TGNAAAAAAAAAAAAAAAAAA 625
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mes 21; Conserv
                                                                               Similarity
                                                             llarity 100.0%; I Conservative 0;
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                                                                              Score 21;
Pred. No.
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                                                               Mismatches
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                                                                              DB 10;
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                                                                                              Length 547;
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US-09-783-590-7037
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 7037
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/783,590 CURRENT FILING DATE: 2000-02-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Dillon, Patrick J. APPLICANT: Haseltine, William A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 08/420,856
PRIOR FILING DATE: 1995-04-12
PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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OTHER INFORMATION: n equals a,t,9, or c
NAME/KEY: misc feature
LOCATION: (109)
OTHER INFORMATION: n equals a,t,9, or c
NAME/KEY: misc feature
LOCATION: (122)
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OTHER INFORMATION: n equals a,t,
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OTHER INFORMATION: n equals a,t,g, or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                           NAME/KEY: misc feature
LOCATION: (409)
OTHER INFORMATION: n equals a,t,g, or
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OTHER INFORMATION: n equals a,t,g, or
                                                              LOCATION: (418)
OTHER INFORMATION: n equals
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                        NAME/KEY: misc feature LOCATION: (428)
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Ruben, Steven M
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US-09-925-301-539
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                                                                                                                                                                                                SOFTWARE: PatentIn Ver. SEQ ID NO 539
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                                                                                                                                                                                                                                                                                                       APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins
FILE REFERENCE: PA106
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 2000-03-08
                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/124,270 PRIOR FILING DATE: 1999-03-12 NUMBER OF SEQ ID NOS: 1694
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OTHER INFORMATION: n equals a,t,g,
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      LOCATION: (1305)
OTHER IFFORMATION: n equals a,t,g, or NAME/KBY: misc_feature
LOCATION: (1344)
                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
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NAME/KEY: misc_feature
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es 21; Conserv
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CURRENT APPLICATION NUMBER: US/09/925/297
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05989
PRIOR ETLING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR ETLING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 928
SOFTWARE: PAtentin Ver. 2.0
SEQ ID NO 171
            APPLICANT: Rosen, Craíg A.

APPLICANT: Ruben, Steven M.

TITLE OF INVENTION: Human Genes, Sequences, an FILE REFERENCE: PO-16.2C1
CURRENT APPLICATION NUMBER: US/09/783,590
CURRENT FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 08/420,856
PRIOR APPLICATION NUMBER: 08/46,731
PRIOR FILING DATE: 1995-04-12
PRIOR FILING DATE: 1994-11-21
NUMBER OF SEO ID NOS: 12485
SOFTWARER: DATE: DATE: 1995-1485
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US-09-925-297-171
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Best Local Similarity 100.0%; P
Watches 21; Conservative 0;
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APPLICANT: Haseltine, Willis
APPLICANT: Li, Haodong
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
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SOFTWARE: PatentIn Ver.
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LOCATION: (1370)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (1400)
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OTHER INFORMATION: n equals a,t,g, or c
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les 21; Conserv
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Pred. No.
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7.6;
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US-09-925-301-759/c
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Sequence 759, Application US/09925301
Patent No. US20020052308A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA106
                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 934
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 419
LENGTH: 60
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                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Pyle, Ruth
APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.552
CURRENT APPLICATION NUMBER: US/09/919,580
CURRENT FILING DATE: 2001-07-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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LENGTH: 58
                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature LOCATION: 54
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LOCATION: (58)
OTHER INFORMATION: n equals a,t,g, or c
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LOCATION: (52)
OTHER INFORMATION: n equals
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LOCATION: (34)
OTHER INFORMATION: n equals a,t,g, or c
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100.0%; Pred. No.
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100.0%; Pred. No
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23;
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CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 759
LENGTH: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; NAME/KEY: misc_feature; LCGATION: (66); OTHER INFORMATION: n equals a,t,g, or c us-09-925-301-759
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US-09-815-343-241/c
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                                                                                                                                  US-09-815-343-241
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 241, Application US/09815343 Patent No. US20010055596A1 GENERAL INFORMATION:
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                      SEQ ID NO 241
                                                                  Matches
                                                                                               Query Match
                                                                                Best Local
                                                                                                                                                                                                                                                                                APPLICANT: Meagher, Madeleine
APPLICANT: Xu, Jiangchun
APPLICANT: Xiu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.504
CURRENT APPLICATION: NUMBER: US/09/815,343
CURRENT FILING DATE: 2001-03-22
NUMBER OF SEQ ID NOS: 1556
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: (65)
OTHER INFORMATION: n equals a,t,g,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
LOCATION: (63)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: n equals a, t, g,
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LOCATION: (6)
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                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (1)...(97)
OTHER INFORMATION: n = A,T,C or G
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                               606 GNAAAAAAAAAAAAAAA 625
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20;
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Pred. No. 23;
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                                                                                Score 20; Pred. No.
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23;
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GENERAL INFORMATION:
APPLICANT: Wang, Ton
APPLICANT: McNeill,
APPLICANT: Watanabe
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US-10-046-935-291/c
                                                                                                   US-10-046-935-291
                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/046,935
CURRENT FILING DATE: 2002-01-15
NUMBER OF SEQ ID NOS: 2239
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 291
LENGTH: 226
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SEQ ID NO 248
LENGTH: 217
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Patent No. US20020099012A1
                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Jiang, Y
APPLICANT: Harlock
APPLICANT: Secrist
APPLICANT: Wang, F
APPLICANT: Stolk,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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Best Local Similarity
Matches 20; Conserv
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Patent No. US20020156011A1
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CURRENT APPLICATION NUMBER: US/09/895,828

CURRENT FILING DATE: 2001-06-29

NUMBER OF SEC.
                                    Matches
                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Henderson, Robert A.
APPLICANT: Kalos, Michael D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER FILE REFERENCE: 210121.527C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                     NAME/KEY: misc_feature
LOCATION: 20, 57, 141, 184, 206
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                        ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: n = A,T,C or
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606 GNAAAAAAAAAAAAAAAA 625
                                                Local Similarity
                                    20;
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Carter, Darrick
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                                                                                                                                                                                                                                                                                                                                                                                   Wang, Aijun
Stolk, John A.
                                                                                                                                                                                                                                                                                                                                                                                                                    Harlocker, Susan
Secrist, Heather
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                                    Conservative
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                                                    Score 20; pred. No.
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Pred. No.
                                      Mismatches
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22;
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                                                                        Length 226;
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21 GNAAAAAAAAAAAAAAAAA

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US-09-834-975-374/c
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; Sequence 46, Application US/10040916
; Patent No. US20020146769A1
; GENERAL INFORMATION:
                                                                                                                           Sequence 374, Application US/09834975 Patent No. US20020110815A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
APPLICANT: Lillie, James
APPLICANT: Brown, Jeffrey
APPLICANT: Bott, Andrew
APPLICANT: Bolt, Andrew
APPLICANT: Van Huffel, Christophe
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE IDENTIFICATION, ASSESSMENT, F
                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
TELECOMMUNICATION. INFORMATION:
TELEPHONE: (617) 498-8224
TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 46:
                                                                                                                                                                                                                                                                                       606 GNAAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                                                                  245 GNAAAAAAAAAAAAAAAA 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/686,878 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/887,029
FILING DATE: 07-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE DESCRIPTION: SEQ ID NO: 46:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute,
STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Jacobs, Kenneth
McCoy, John
LaVallie, Edward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/040,916
FILING DATE: 07-Jan-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 265 base pairs
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                                                                                                                                                                                                                                                                                                                                                Conservative
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Evans, Cheryl
Spaulding, Vikki
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100.0%; Pred. No.
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    PREVENTION, AND THERAPY
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US-09-783-590-10830
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; Patent No. US20020110850A1
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US-09-777-564-625/c
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; NAME/KEY: misc_feature
; LOCATION: (1)...(291)
; OTHER INFORMATION: n = A,T,C or
US-09-834-975-374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Algate, Paul A.
APPLICANT: Mannion, Jane
APPLICANT: Mannion, Jane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/777,564
CURRENT FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 1730
SOFTWARE: FastSEQ for Window Version 4.0
SEQ ID NO 625
LENGTH: 451
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 374
LENGTH: 291
TYPE: DNA
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Patent No. US20020022591A1
                  APPLICANT: Dillon, Patrick J.
APPLICANT: Haseltine, William A.
APPLICANT: Li, Haodong
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2
FILE REFERENCE: PO-16.2C1
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CURRENT APPLICATION NUMBER: US/09/783,590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER FILE REFERENCE: 210121.493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/197,538 PRIOR FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/834,975
CURRENT FILING DATE: 2001-04-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: n = A,T,C or G
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                                                                                                                                                                                                                                                                                                     373 GNAAAAAAAAAAAAAAAA 354
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                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.2%; Score 20;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                    3.2%; Score 20; DB 100.0%; Pred. No. 21;
                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                       DB 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 10830
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PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT FILING DATE: 2000-02-15 PRIOR APPLICATION NUMBER: 08/420,856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 12485
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LOCATION: (8)
OTHER INFORMATION: n equals a,t,g, or c
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                                                                                                                                                                                                                                                                    LOCATION: (357)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc feature LOCATION: (318)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc feature LOCATION: (310)
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OTHER INFORMATION: n equals a,t,g,
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OTHER INFORMATION: n equals a,t,g,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                              LOCATION: (396)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                       OTHER INFORMATION: n equals a,t,g, or NAME/KEY: misc feature
                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: n equals a,t,g, or
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                   OTHER INFORMATION: n equals a,t,g, or c
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                                                                                                                                                                                                                                                                                                           NAME/KEY: misc feature
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NAME/KEY:
                                    NAME/KEY: misc feature LOCATION: (404)
                                                                                                                                                                                 NAME/KEY: misc feature
                                                                                                                                                                                                                           LOCATION: (364)
                                                                                                                                                                                                                                              NAME/KEY: misc feature
  misc feature
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CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTMARE: Patentin Ver. 2.0
SEQ ID NO 819
LENGTH: 648
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies FILE REFERENCE: PA106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Rosen et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
LOCATION: (419)
OTHER INFORMATION: n equals a,t,9,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc feature LOCATION: (493)
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OTHER INFORMATION: n
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FEATURE:
NAME/KEY: misc_feature
LOCATION: (369)
OTHER INFORMATION: n equals a,t,9, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: n equals a,t,g, or
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LOCATION: (518)
OTHER INFORMATION: n equals
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             OTHER INFORMATION: n equals a,t,g,
NAME/KEY: misc_feature
LOCATION: (584)
                                                                   OTHER INFORMATION: n equals a,t,g,
NAME/KEY: misc_feature
LOCATION: (565)
                                                                                                                         OTHER INFORMATION: n equals a,t,g, NAME/KEY: misc_feature LOCATION: (547)
                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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OTHER INFORMATION: n equals a,t,g, or
                                                                                                                                                                                 NAME/KEY: misc_feature LOCATION: (544)
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[00.0%; Pred. No.
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APPLICANT: Wang, Aijun

APPLICANT: Skeiky, Yasii A.W.

APPLICANT: Hepler, William

TITLE OF INVENTION: COMPOSITIONS AND METHODS

TITLE OF INVENTION: COMPOSITIONS AND METHODS

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CA

FILLE REFERENCE: 21021, 427C23

CURRENT APPLICATION NUMBER: US/09/759,143

CURRENT FILING DATE: 2001-01-12

NUMBER OF SEQ ID NOS: 934

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 32

LENGTH: 789

TYPE: DNA

ORGANISM: Homo sapien
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                                                                                      GENERAL INFORMATION:
                                                                                                     Sequence 32, Application US/09780669 Patent No. US20020051977A1
                  APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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Patent No. US20020022248A1
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LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
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mes 20; Conserv
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                                    Xu, Jiangchun
Dillon, Davin C.
Mitcham, Jennifer L.
Jiang, Yuqui
Henderson, Robert A.
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Vedvick,
                              Harlocker, Susan L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Li, Samuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Carter, Darrick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stolk, John A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jiang, Yuqui
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Retter, Marc W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fanger, Gary R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kalos, Michael D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Henderson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Harlocker,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dillon, Davin C.
                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Craig H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jiangchun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPOSITIONS AND METHODS FOR THE THERAPY AND DIAGNOSIS OF PROSTATE CANCER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Susan L.
                                                                                                                                                                                                                                                                3.2%; Score 20;
100.0%; Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                           10; Length 789;
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RESULT 34
US-09-030-606-32/c
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                                                                                                                                                                                                                                                                                                                                                                                      Patent No. ÚSZÕÒ20081580A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Xi, Jiangchun
APPLICANT: Dillon, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNODIAGNOSIS OF PROSTATE CANCER AND METHOD
NUMBER OF SEQUENCES: 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 943
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 32, Application US/09030606
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INFORMATION FOR SEQ ID NO:
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APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND FILE REFERENCE: 210121.427C24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                     ATTORNEY AGENT INFORMATION:
NAME: Maki, David J.
NEGISTRATION NUMBER: 31,392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                           REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature LOCATION: (1)...(789)
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                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center,
                                                                                                                                                                                                                                                                                                                  STREET: 6300 C
CITY: Seattle
STATE: WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   606 GNAAAAAAAAAAAAAAAA 625
                                                                                                                                                APPLICATION NUMBER: US/0 FILING DATE: 25-FEB-1998
                   TELEFAX:
                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                COUNTRY:
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les 20; Conserv
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Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
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Hepler, William
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Carter, Darrick
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                   (206)
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                 682-6031
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                                                                210121.428C3
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                                                                                                                                                                                                                                                                                                                                              701 Fifth Avenue
                                                                                                                                                                                             Version #1.30
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US-09-822-827-32/c
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US-09-115-453-32/c
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US-09-115-453-32
                                                                                                                                                                                                                                                                                                                  Sequence 32, Application US/09115453B Patent No. US20020090372A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
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Patent No. US20020081680A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
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                                                                                                                                        NUMBER OF SEQ ID NOS: 228
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER ETILE REFERENCE: 210121.534C1
                                                                                                                                                                                            TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND TITLE OF INVENTION: METHODS FOR THEIR USE FILE REFERENCE: 210121.427C4
CURRENT APPLICATION NUMBER: US/09/115,453B
CURRENT FILING DATE: 1998-07-14
                                                                                                                                                                                                                                                                                  APPLICANT: Xu, Jiangchun APPLICANT: Dillon, Davin C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
         NAME/KEY: misc_feature
LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
                                                                     LENGTH: 789
TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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LENGTH: 789 base pairs
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Pred. No.
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US-09-783-590-8086/c
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Best Local Similarity
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                                                                                                                                                        SOFTWARE: PatentIn Ver. SEQ ID NO 8086
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CURRENT FILING DATE: 2001-09-12
PRIOR APPLICATION NUMBER: 60/231,572
PRIOR FILING DATE: 2000-09-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS, TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUM TITLE OF INVENTION: AND USES THEREOF FILE REFERENCE: CL000817
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Dillon, Patrick J.
APPLICANT: Haseltine, William A.
APPLICANT: Li, Haodong
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: YAN, Chunhua et al
                                                                                                                                                                                           CURRENT FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 08/420,856
PRIOR FILING DATE: 1995-04-12
PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
NUMBER OF SEQ ID NOS: 12485
                                                                                                                                                                                                                                                                                                                                  APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences,
FILE REFERENCE: PO-16.2C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/783,590 CURRENT FILING DATE: 2000-02-15
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              FEATURE: misc feature
NAME/KEY: misc feature
LOCATION: (10)
OTHER INFORMATION: n equals a,t,g, or c
                                                                                             LENGTH: 51
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: n = A,T,C or
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NAME/KEY: misc feature
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100.0%; Pred. No.
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. 20;
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                                                                                                                                                                                                                                                                                                                                                               and Expression Products 16.2
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US-09-919-580-705/c
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                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/919,580
CURRENT FILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 934
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 705
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TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2
FILE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2
FILE OF INVENTION (1997)
FILE OF THE OFFICE OF OFFICE OF OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFFICE OFF
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APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
FITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.552
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                                                                 ORGANISM: Homo sapiens
                                                                                                         TYPE: DNA
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ORGANISM: Homo sapiens
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NAME/KEY: misc_feature
                                                                                                                                             LENGTH: 53
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OTHER INFORMATION: n equals a,t,g, or c
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APPLICANT: Haseltine, William A.
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OTHER INFORMATION: n equals a,t,g, or c
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Ruben, Steven M.
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100.0%; Pred. No.
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100.0%; Pred. No.
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CURRENT APPLICATION NUMBER: US/09/783,590
CURRENT EILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 08/420,856
PRIOR FILING DATE: 1995-04-12
PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
NUMBER OF SEQ ID NOS: 12485
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 3220
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US-09-783-590-3220/c
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                NUMBER OF SEQ ID NOS: 934

SOFTWARE: FRASTSEQ for Windows Version 4.0
SEQ ID NO 171
LENGTH: 56
                                                                                                        APPLICANT: Pyle, Ruth
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.552
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                                                              CURRENT APPLICATION NUMBER: US/09/919,580 CURRENT FILING DATE: 2001-07-30 NUMBER OF SEQ ID NOS: 934
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APPLICANT: Dillon, Patrick J.
APPLICANT: Haseltine, Willia
APPLICANT: Li, Haodong
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TYPE: DNA
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APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2
FILE REFERENCE: PO-16.2C1
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OTHER INFORMATION: n equals a,t,g, or c
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ORGANISM: Homo sapiens
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OTHER INFORMATION:
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Local Similarity 100.0%; F
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100.0%; Pred. No.
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US-09-919-580-713/c
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 677
LENGTH: 56
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Best Local :
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APPLICANT: Xu, Jlangchun
APPLICANT: Secrist, Heather
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.552
CURRENT APPLICATION NUMBER: US/09/919,580
CURRENT FILING DATE: 2001-07-30
                                                                                                             NUMBER OF SEQ ID NOS: 934
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 713
LENGTH: 56
                                                                                                                                                                                            APPLICANT: Pyle, Ruth
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.552
CURRENT APPLICATION NUMBER: US/09/919,580
CURRENT FILING DATE: 2001-07-30
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NAME/KRY: mlsc_feature
LOCATION: 49
OTHER INFORMATION: n = A,T,C or G
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LOCATION: 53
OTHER INFORMATION: n = A,T,C or G
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TYPE: DNA
OGGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: 35, 37, 42, 56
OTHER INFORMATION: n = A,T,C or G
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US-09-925-301-758/c
                                                                          В
                                                                                                                                                                                                             US-09-925-301-758
Search completed: November 8, 2002, 02:06:25 Job time : 100.162 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. US20020052308A1
GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies FILE REFERENCE: PA106
                                                                                                                                              Matches
                                                                                                                                                               Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Rosen et al
                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
LOCATION: (45)
OTHER INFORMATION: n equals a,t,9, or
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION: (36)
OTHER INFORMATION: n equals a,t,9,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: (38)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                                                                                                                                OTHER INFORMATION: n equals a,t,g, or c
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LOCATION: (40)
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                                                                                                                                                                 100.0%;
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100.0%; Pred. No.
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Or
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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Gapop 60.0 , Gapext 60.0
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Copyright (c) 1993 - 2002 Compugen Ltd.
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/cgn2_6/ptodata/1/pna/US102B<u>-</u>COMB
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11225.881 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

19 20 21	11 12 14 16	c 109876554321	Result No.
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US-09-652-109-9370 US-09-652-121-6191 US-09-652-128-7518	PCT-US01-18569-111 US-10-264-049-111 US-10-108-260A-197 US-10-208-408-26 US-10-295-262-26 US-60-308-868-26 US-60-371-168-7188	PCT-US01-45395-210 US-09-825-294-210 US-09-877-701-10876 US-09-970-966-210 US-10-212-677-210 PCT-US01-45395-214 US-09-970-966-214 US-10-212-677-214 US-10-212-677-214 US-10-212-677-214	ID
Sequence 9077, Ap Sequence 9370, Ap Sequence 6191, Ap Sequence 7518, Ap	1111 26 26 271	Sequence 210, App Sequence 210, App Sequence 210, App Sequence 211, App Sequence 214, App Sequence 214, App Sequence 214, App Sequence 214, App Sequence 214, App Sequence 214, App Sequence 214, App Sequence 214, App Sequence 9, Appli	Description

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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: CORIXA CORPORATION

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Hing, Steven P.

APPLICANT: Molesh, David Alan

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.48401PC

CURRENT APPLICATION NUMBER: PCT/US01/45395

CURRENT FILING DATE: 2001-11-13

NUMBER OF SEQ ID NOS: 215

SOPTWARE: FRSSTSEQ for Windows Version 4.0

SEQ ID NO 210

LENGTH: 625

TYPE: NAME
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                                     CAGCGACAGGCGCACACCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
                                                                                                                                          GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                                                                         AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
                                                                                                                      GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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APPLICANT: Xu, Jiangchun

APPLICANT: Xtolk, John A.

APPLICANT: Algate, Paul A.

APPLICANT: Algate, Paul A.

APPLICANT: Filng, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA

FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 210

LENGTH: 625
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Pred. No. 6.3e-125;
Mismatches 0;
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ORGANISM: Homo sapiens

TYPE: DNA

Query Match Best Local Similarity Matches 625; Conserv

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Sequence 10876, Application US/09867701

Sequence 10876, Application US/09867701

APPLICANT: Adjate, Paul A.

APPLICANT: Jones, Robert

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEQ ID NOS: 10912

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 10876

LENGTH: 625

TYPE: DNA

ORGANISM: Homo sapiens

FEATURE:

NAME XEVY: TASC foature
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                                                                                                                                                                        Score 625; DB 33; Pred. No. 6.3e-125;
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ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: 607
OTHER INFORMATION: n = A
US-09-970-966-210
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.484C6

CURRENT APPLICATION NUMBER: US/09/970,966

CURRENT FILING DATE: 2001-10-02

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 210

LENGTH: 625

TYPE: DNA

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Best Local Similarity
Matches 625; Conserv
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APPLICANT: Molesh, David Ali
APPLICANT: Fling, Steven P.
APPLICANT: Xu, Jiangchun
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                                                                                                                                  Conservative
                                                                                                                                         100.0%; Score 625; DB 36; 100.0%; Pred. No. 6.3e-125;
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SOFTWARE: FastSEQ for
SEQ ID NO 214
LENGTH: 1897
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                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                      Matches 606;
                                                                                                                                                                                                                                                                                         APPLICANT: Algate, Paul A.
APPLICANT: #1ing, Steven P.
APPLICANT: Molesh, David Alan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 2101.21.48401PC
CURRENT APPLICATION NUMBER: PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 215
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                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
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ilarity 100.0%;
Conservative
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                                                                                                                                                                Score 606; DB 1; L
Pred. No. 5.3e-121;
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Y 1 AGTTCTCCTTGCAGAGGACTGGCGCGCGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60	Db Qq
Query Match 100.0%; Score 625; DB 42; Length 625; Best Local Similarity 100.0%; Pred. No. 6.3e-125; Matches 625; Conservative 0; Mismatches 0; Indels 0; Gaps	M B Q
OTHER INFORMATION: n = A,T,C or G S-10-212-677-210	Š
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NUMBER OF SEQ ID NOS: 288  SOFTWARE: FastSEQ for Windows Version 4.0  SEO ID NO 210	'n
PLICATION NO	
NVENTION: AND DIAGNOSIS OF OVARIAN CANCER ENCE: 210121 484C7	
MCNeill, Patricia D. MCNeill, Patricia D. MCNEITION: COMPOSITIONS AND METHODS FO	
APPLICANT: Xu, Jiangchun APPLICANT: Fanger, Gary R. ADDITOANT: Harlocker Susan I	
GENERAL INFORMATION: APPLICANT: Chenault, Ruth A.	
SULT 5 -10-212-677 Sequence 21	RESU US-1
b 601 ACAGTGNAAAAAAAAAAAAAAAA 625	밁
601	Q
b 541 AACACTGTTTCAAACCCGGGGAGTTGGCTGTGTTAAAGAAAG	DЬ
541 AAC	Оу
b 481 GGTACCAATTTTTCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAG	DЬ
481 GGTACCAATTTTTCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAG	Qy
b 421 TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480	B
421	Qy
561 TGGGCAGACACGAGCAGGAGAGAGAGAAAAGAAAAACACACGCATGAGAACACAG 420	D
361 TGGGCAGACACGAGCAGGAGGGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACAC	Q
b 301 CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC 360	망
301 CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC	Qy
241 GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 3	Db
241	Qγ
b 181 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 240	DЪ
181	νQ

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:, TYPE: DNA
: ORGANISM: Homo sapiens
: FEATURE:
: NAME/KEY: misc_feature
: LOCATION: (1)...(1897)
: OTHER INFORMATION: n = A,T,C or
US-09-825-294-214
                                                                                                                                                                                                                                                                                          APPLICANT: Xu, Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Fling, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 214

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Best Local :
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121 CAGCGACAGGCGGCAGCACCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC
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                                                                                                                                                        Local Similarity
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                          GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGCGCTTCTCGTGGTTGGCGTGCTG
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                                                                                                                                                    97.0%; Score 606; DB 31; 100.0%; Pred. No. 5.3e-121;
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APPLICANT: Molesh, David Alan
APPLICANT: Molesh, David Alan
APPLICANT: Filing, Steven P.
APPLICANT: YL, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.48406
CURRENT APPLICATION NUMBER: US/09/970,966
CURRENT APPLICATION NUMBER: US/09/970,966
CURRENT FILING DATE: 2001-10-02
NUMBER OF SEQ ID NOS: 215
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 214
LENGTH: 1897
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; ORGANISM: Homo sapiens
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Best Local Similarity
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                                                                                                                                                              97.0%; Score 606; DB 36; ilarity 100.0%; Pred. No. 5.3e-121; Conservative 0; Mismatches 0;
                                                                                                                                                                                          Length 1897;
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NAME/KEY: misc_feature

; LOCATION: (1)...(43729)

; OTHER INFORMATION: n = A,T,C

US-60-261-974-9
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TITLE OF INVENTION: I
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                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 606; Conserv
                                                                                                                                                                                                                                                                                                                        SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/60/261,974
CURRENT FILING DATE: 2001-01-17
NUMBER OF SEQ ID NOS: 237
                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                    SOFTWARE:
                                                                                                                                       41523
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 41403 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 41344
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                                                                                                                                                                                97.0%; Score 606; DB 70;
100.0%; Pred. No. 2.2e-121;
Vative 0; Mismatches 0;
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S S S S S S S S S S S S S S S S S S S	ORCE OF STREET	214 , Application US/10212677 )RMATION: Chenault, Ruth A. Xu, Jiangchun Fanger, Gary R. Harlocker, Susan L. MCNeill, Patricia D. MCNEILL, PATRICIA D. WENTION: COMPOSITIONS AND METHODS FOR THE THE AVENTION: AND DIAGNOSIS OF OVARIAN CANCER ENCE: 210121.484C7 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,677 JLICATION NUMBER: US/10/212,6	
X B O	ery M st Lc tches	atch 97.0%; Score 606; DB 42; Length 1897; cal Similarity 100.0%; Pred. No. 5.3e-121; 606; Conservative 0; Mismatches 0; Indels 0; Gap	
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₹	181	ACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGG	_

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TITLE OF INVENTION: NUCLEAC ACIDS, PROTEINS AND FILE REFERENCE: PA133PCT CURRENT APPLICATION NUMBER: PCT/USO1/18569 CURRENT FILING DATE: 2001-06-07 PRIOR APPLICATION NUMBER: 60/209,467 PRIOR FILING DATE: 2000-06-07 NUMBER OF SEQ ID NOS: 4360 SOFTWARE: PATENTIN VET. 2.0 SEQ ID NO 111 SEQ ID NO 111 LENGTH: 1608
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                                                                                                                       Matches 605;
                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 111, Application PC/TUS0118569
GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc.
                                                                                                                                                                                                                                                                         LOCATION: (18)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: SITE
LOCATION: (22)
OTHER INFORMATION: n equals a,t,g, or c
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LOCATION: (32)
OTHER INFORMATION: n
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LOCATION: (87)
                                                                                                                                                                                            OTHER INFORMATION: n
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                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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                                                                                                                                      Local Similarity
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GCCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 1044
                                                       AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 984
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                                                                                                                              Score 555; DB 1;
Pred. No. 4.6e-110;
                                                                                                                    Mismatches
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GENERAL INFORMATION:
APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
                                                                                                                                                                                                                                                                                      SEQ ID NO 111
                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: PCT/US01/18569
PRIOR FILING DATE: 2001-06-07
PRIOR APPLICATION NUMBER: US 60/209,467
PRIOR ETILING DATE: 2000-06-07
NUMBER OF SEQ ID NOS: 4360
SOFTWARE: Patentin Ver. 3.1
                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/264,049 CURRENT FILING DATE: 2002-10-04
                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: PA133P1
                                                                                                                                                                                                                                                        LENGTH: 16
NAME/KEY: misc_feature
LOCATION: (87)..(87)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                  NAME/KEY: misc_feature
LOCATION: (22). (22)
OTHER INFORMATION: n equals
                                                                           NAME/KEY: misc_feature LOCATION: (32)..(32)
                                                                                                                                                                                            FEATURE:
NAME/KEY: misc_feature
LOCATION: (18)...(18)
                                                            OTHER INFORMATION: n equals
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                                                                                                                                                                               OTHER INFORMATION: n equals
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                                                  FEATURE:
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Sequence 197, Application US/10108260A
GENERAL INFORMATION:
APPLICANT: HELIX RESEARCH INSTITUTE
FITLE OF INVENTION: NOVEL full length cDNA
FILE REFERENCE: H1-A0106
CURRENT FILING DATE: 2002-03-27
NUMBER OF SEQ ID NOS: 5458
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 197
LENGTH: 1832
TYPE: DNA
OCOLNICAL INFORMATION
                                                                                       ; ORGANISM: Homo sapiens US-10-108-260A-197
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Best Local Similarity
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                            Score 555; DB 40;
Pred. No. 4.4e-110;
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                                                                                                                        US-10-208-408-26
                                                                                                                                                                                                                                                                                                   Sequence 26, Application US/10208408
GENERAL INFORMATION:
APPLICANT: Schebye, Xiao Min
TITLE OF INVENTION: GENES REGULATED BY PEROXISOME PROLIFERATOR-ACTIVATED RECEPTOR
FILE REFERENCE: PA-0048-1 US
CURRENT APPLICATION NUMBER: US/10/208,408
CURRENT FILING DATE: 2002-07-29
CURRENT FILING DATE: 2002-07-29
                                                                                                                                                                                                 SOFTWARE: PERL Program
SEQ ID NO 26
LENGTH: 1866
TYPE: DNA
                                                                            Query Match
Best Local
                                                               Matches
                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/308,868
PRIOR FILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 55
                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                      OTHER INFORMATION: Incyte ID
                                                                                                                                                     NAME/KEY: misc_feature
                                                                                                                                                                    FEATURE:
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                                                                          Local Similarity
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                                                               Conservative
                                                                             99.8%;
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                                                                            Score 555; DB 42;
Pred. No. 4.4e-110;
                                                               Mismatches
                                                                                                                                         227484.13
                                                                                             Length 1866;
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Sequence 26, Application US/60295262
(GENERAL INFORMATION:
GENES REGULATED BY PEROX
TITLE OF INVENTION: GENES REGULATED BY PEROX
TITLE OF INVENTION: GAMMA AGONIST
FILE REFERENCE: PA-0048 P
CURRENT APPLICATION NUMBER: US/60/295,262
CURRENT FILING DATE: 2001-06-01
NUMBER OF SEQ ID NOS: 51
SOFTWARE: PERL PROGram
SEQ ID NO 26
LENGTH: 1866
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                                                             S
                                                                                                                                     ; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte
US-60-295-262-26
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                                                                                        Matches
                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                   1141 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGGCAACGGGCGCTGCACAAAGCG 1200
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                                                                                        605;
                                                                                                  Similarity
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                                                                                     Conservative
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                                                                                                                                               No:
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                                                                                               Score 555; DB 73; Pred. No. 4.4e-110;
                                                                                    Mismatches
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                                                                                                         Length 1866;
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RESULT 16
US-60-308-868-26
Sequence 26, Application US/60308868
GENERAL INFORMATION:
APPLICANT: Schebye, Xiao Min
TITLE OF INVENTION: GENES REGULATED BY PEROXISOME PROLIFERATOR-ACTIVATED RECEPTOR
FILE REFERENCE: PA-0048-1 P
CURRENT APPLICATION NUMBER: US/60/308,868
CURRENT FILING DATE: 2001-07-30
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SOFTWARE: PERL Program
SEQ ID NO 26
LENGTH: 1866
TYPE: This
                                                                                                                                                      Matches 605;
                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                   NAME/KEY: misc_feature
OTHER INFORMATION: Incyte
                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                             TYPE: DNA
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                                                                                                          1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCGCTGCACAAGCG
              CAGCGACAGGCGCAGCACCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
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Conservative
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                                                                                                                                                                                                                      No:
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                                                                                                                                                              Score 555; DB 74;
Pred. No. 4.4e-110;
                                                                                                                                                   Mismatches
                                                                                                                                                                           Length 1866;
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US-09-644-873-9077
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SEQ ID NO 9077
LENGTH: 1917
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 9077, Application US/09644873 GENERAL INFORMATION:
                                                                                                                                                                                     Matches 605;
                                                                                                                                                                                                                                                                                              APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: Holtzman, Douglas A.
TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND
TITLE OF INVENTION: THEREFOR
TITLE OF INVENTION: THEREFOR
CURRENT FILING DATE: 2000-08-28
PRIOR APPLICATION NUMBER: 05/05/644,873
CURRENT FILING DATE: 2000-08-28
PRIOR APPLICATION NUMBER: 60/151,064
PRIOR FILING DATE: 1999-08-27
NUMBER OF SEQ ID NOS: 11286
TOTAL TO THE SEQ ID NOS: 11286
                                                                                                                                                                                               Query Match
Best Local Similarity
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                                                                                                     61 GGCGCTGTCGGTGGAGGGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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                       CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                          CAGCGACAGGCGGCAGCACCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
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                                                                                                                                                                                   Score 555; DB 25;
Pred. No. 4.4e-110;
0; Mismatches 1;
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CAGCGACAGGCGGCAGCACAGCACCTGCACGACACACCCGCCGAAACTGCTGCGAGGACAC 180	GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120	AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60	<pre>latch 88.8%; Score 555; DB 17; Length 1917; local Similarity 99.8%; Pred. No. 4.4e-110; local Similarity 99.8%; Pred. No. 1.4e-110; local Similarity 99.8%; Gaps 1; Indels 0; Gaps</pre>	ULT 17  09-371-168-7188 equence 7188, Application US/09371168 ENERAL INFORMATION: APPLICANT: Glucksmann, M. Alexandra APPLICANT: Glucksmann, M. Alexandra TITLE OF INVENTION: NUCLEIC ACID MOLECULES DERIVED FROM A TITLE OF INVENTION: HUMAN BRAIN LIBRARY FILE REFERENCE: MLN98-37PA CURRENT APPLICATION NUMBER: US/09/371,168 CURRENT FILING DATE: 1999-08-10 EARLIER APPLICATION NUMBER: 60/095,907 EARLIER APPLICATION NUMBER: 60/095,907 EARLIER FILING DATE: 1998-08-10 EARLIER FILING DATE: 1998-08-10 EARLIER FILING DATE: 1998-0-05 NUMBER OF SEQ ID NOS: 8285 SOFTWARE: FastSEQ for Windows Version 3.0 EQ ID NO 7188 ENGTH: 1917 TYPE: DNA ORGANISM: Homo sapiens 09-371-168-7188	ACAGTG 1746	ACAGTG 606	AACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAG	GGTACCAATTTTTCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAG	TARATGAATAAACCATAAAATATTTAGCCCCTCTGTTCTGT	TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAGAAA	CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC 360	GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 300	CCTGTALAGGACCGGTTGATGACCGACCTGACGTAGAAAACGTCTCCCGAGAAGGGGAG 240	AGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC 13

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US-09-652-109-9370

Sequence 9370, Application US/09652109

GENERAL INFORMATION:

APPLICANT: MCCARTHY, Sean A.

TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES

TITLE OF INVENTION: THEREFOR

FILE REFERENCE: 1600.1180-001

CURRENT APPLICATION NUMBER: US/09/652,109

CURRENT FILING DATE: 2000-08-30

PRIOR APPLICATION UNMBER: 60/151,128

PRIOR FILING DATE: 1999-08-30

NUMBER OF SEQ ID NOS: 10105

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 9370

LENGTH: 1917

TYPE: DNA

TYPE: DNA

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           GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 300
                                                            CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 240
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Pred. No. 4.4e-110;
0; Mismatches 1;
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CURRENT APPLICATION NUMBER: US/09/652,121
CURRENT ELING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/151,129
PRIOR FILING DATE: 1999-08-30
NUMBER OF SEQ ID NOS: 7615
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6191
LENGTH: 1917
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Sequence 6191, Application US/09652121
GENERAL INFORMATION:
APPLICANT: Distefano, Peter
TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
TITLE OF INVENTION: THEREFOR
FILE REFERENCE: 1600.1188-001
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TYPE: DNA
ORGANISM: Homo sapiens
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Qy 421 TAAATGAATAAAACCATAAAATATTTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480	QY 361 TGGGCAGACACGAGCAGGAGGAGGAGAGAGAAAAGAAAA	QY 301 CATGATCCTCCGAATCTGGTTGGGCATCAGCATACGGCCAATGTCACAACAATCAGCCC 360	OY 241 GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGCTTTGGCCGCAGC 300	QY 181 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 240	QY 121 CAGCGACAGGCGGCAGCACCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC 180	QY 61 GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120	QY 1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60	Query Match 88.8%; Score 555; DB 25; Length 1917; Best Local Similarity 99.8%; Pred. No. 4.4e-110; Matches 605; Conservative 0; Mismatches 1; Indels 0; Gaps 0	Qy 421 TANATGANTANAACCATAAAATATTTACCCCTTGTTTGTTCTTGTCTTACTGACAAATATATAAATAA
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Qy 361	Qy 301 Db 1586	Qy 241 Db 1526	Qy 181 Db 1466	Qy 121 Db 1406	Qy 61 Db 1346	Qy 1 Db 1286	Query Ma Best Loc Matches	RESULT 21 US-09-652-128 Sequence 75 GENERAL INF GENERAL INF GENERAL INF GENERAL INF INTLE OF I FILE REFER CURRENT FI CURRENT FI PRIOR APPL PRIOR APPL INTERIOR OF I NUMBER OF SOFTWARE: SEQ ID NO 7 INTERIOR IDNA ORGANISM: US-09-652-128	Qy 601 Db 1886	Oy 541 Db 1826	Qy 481 Db 1766	Qy 421 Db 1706	Qy 361 Db 1646
TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAGAAA	CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC 3	GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 3	CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 2	CAGCGACAGGCGGCACCACACCACCACGAACACCCGCCGAAACTGCTGCGAGGACAC 1	GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 1	AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCCAACGGGCGCTGCACAAAGCG 6	atch 88.8%; Score 555; DB 25; Length 1917; cal Similarity 99.8%; Pred. No. 4.4e-110; caps 605; Conservative 0; Mismatches 1; Indels 0; Gaps	F 21 652-128-7518 IERCE 7518, Application US/09652128 ERAL INFORMATION: LICANT: HOLTZMAN, Andrew W. PLICANT: HOLTZMAN, DOUGlas A. FLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES FLE OF INVENTION: NOWEL NUCLEIC ACID MOLECULES AND USES FLE REFERENCE: 1600.1171-001 ER RENT APPLICATION NUMBER: US/09/652,128 REENT APPLICATION NUMBER: 05/151,133 FOR FILING DATE: 2000-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE: 1999-08-30 FOR FILING DATE:	ACAGTG 606         ACAGTG 1891	AACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAG	GGTACCAATTTTCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAG	TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 4	TGGGCAGACACGAGCAGGAGGAGGAGAGAGAGAAAAGAAAAACACAGGATGAGAACACAG 4
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US-09-652-914-8473
Sequence 8473, Application US/09652914
GENERAL INFORMATION:
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-652-914-8473
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SEQ ID NO 8473
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TITLE OF INVENTION: HOVEL NUCLEIC ACID M.
TITLE OF INVENTION: THEREFOR
FILE REFERENCE: 1600.1193-001
CURRENT APPLICATION NUMBER: US/09/652,914
CURRENT FILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 60/152,112
PRIOR FILING DATE: 1999-08-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID
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                                                                                           CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAACGTCTCCGAGAAGGGGAG
                                                                             CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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; Sequence 2477, Application US/09652917
; GENERAL INFORMATION:
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SEQ ID NO 2477
LENGTH: 1917
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CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/151,422
PRIOR FILING DATE: 1999-08-30
NUMBER OF SEQ ID NOS: 3855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Shyjan, Andrew W. APPLICANT: Holtzman, Douglas A. APPLICANT: Distefano, Peter TITLE OF INVENTION: NOVEL NUCLEIC TITLE OF INVENTION: THEREFOR FILE REFERENCE: 1600.11170-001
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               CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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                                                                                      TARATGAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480
                                                                                                                    CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
TGGGCAGACACGAGCAGGGGGGGGAGAGACAGAGAAAAGAAAAACACAGCATGAGAACACAG 420
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RESULT 26
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; SEQ ID NO 4454
; LENGTH: 1918
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PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: 60/106,443
PRIOR FILING DATE: 1998-10-30
PRIOR APPLICATION NUMBER: 60/107,257
PRIOR FILING DATE: 1998-11-05
PRIOR APPLICATION NUMBER: 60/126,906
PRIOR APPLICATION NUMBER: 60/126,906
PRIOR APPLICATION NUMBER: 60/126,906
PRIOR FILING DATE: 1999-03-30
PRIOR FILING DATE: 1999-03-30
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Best Local Similarity
Matches 605; Conserv
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APPLICANT: HOLLZMAN, Douglas A.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES DERIVED FROM A
TITLE OF INVENTION: HUMAN FETAL SKIN LIBRARY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/397,022
CURRENT FILING DATE: 1999-09-15
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PRIOR FILING DATE: 1999-04-30
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1707 TAAATAAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 1766
                                                                         1647 TGGGCAGACACGAGCAGGAGGGAGAGACAGAGAAAAGAAAAACACAGCATGAGAACACAG
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                 421 TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480
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Pred. No. 4.4e-110;
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Qy	541 AACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAAGACCATTAAATGCTTTAG 600	
ДЪ	actgtttcaaacccggggagttggctgtgttaaagaaaga	
Qy	601 ACAGTG 606	
Дb	1886 ACAGTG 1891	
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	andra ID MOLEC	
	CURRENT APPLICATION NUMBER: US/09/801,833 CURRENT FILING DATE: 2001-03-13 PRIOR APPLICATION NUMBER: 09/371,168 PRIOR FILING DATE: 1999-08-10	
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ω	PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1998-11 NUMBER OF SEQ ID NOS: 828 SOFTWARE: FastSEQ for Win EQ ID NO 7188 LENGTH: 1917	
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3 to C	lest Local Similarity 99.8%; Pred. No. 4.4e-110; lest Local Similarity 99.8%; Pred. No. 4.4e-110; local Similarity 60; Gaps	
g 9	1 AGTTCTCCTTGCAGAGGACTGGCGCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60	
Ş	GCGCTGTCGGTGGAGTGCGCATGTACGCGCAAGGCGCTTCTCGTGGT	
DЬ	GCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGC	
õ	121 CAGCGACAGGCGGCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC 180	
ф	GACAGGCGGCAGCACACCTGCACGAACACCCGCCGAAACTGCTGCGAGGAC	
Оy	181 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAAGGGGAG 240	
Qy	– വ	
망	AGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAG	
Qy	301 CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC 360	
Вb	ATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC	
Qy	361 TGGGCAGACCAGCAGGAGGAGAGAGAGAAAAGAAAAACACAGCATGAGAACACAG 420	
망	CAGACACGAGCAGGAGGGAGAGACAGAGAAAAGAAAAAACACAGCATGAGAACAC	
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US-09-808-383-4454
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PRIOR ETLING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: US 60/100,465
PRIOR ETLING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: US 60/106,443
PRIOR FILING DATE: 1998-10-30
PRIOR APPLICATION NUMBER: US 60/107,257
PRIOR FILING DATE: 1998-11-15
PRIOR APPLICATION NUMBER: US 60/126,906
PRIOR APPLICATION NUMBER: US 60/126,906
PRIOR APPLICATION NUMBER: US 60/132,099
PRIOR APPLICATION NUMBER: US 60/132,099
PRIOR FILING DATE: 1999-03-30
PRIOR APPLICATION NUMBER: US 60/132,099
PRIOR FILING DATE: 1999-04-30
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4454
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CURRENT APPLICATION NUMBER: US/09/808,383
CURRENT FILING DATE: 2001-03-13
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TITLE OF INVENTION: NUCLEIC ACID MOLECULES DERIVED
TITLE OF INVENTION: HUMAN FETAL SKIN LIBRARY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Gearing, David P. APPLICANT: Holtzman, Dougla
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3TH: 1918
CAGCGACAGGCGCAGCACAGCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC
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                                               GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
                                                              GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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Pred. No. 4.4e-110;
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; LENGTH: 1925
; TYPE: DNA
; ORGANISM: Homo s
US-09-652-121-7293
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SOFTWARE: Fast
SEQ ID NO 7293
FENGTH: 1925
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Best Local
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TITLE OF INVENTION: NOVEL NUCLEIC ACID MOTITULE OF INVENTION: THEREFOR
FILE REFERENCE: 1600.1188-001
CURRENT APPLICATION NUMBER: US/09/652,121
CURRENT ETLING DATE: 2000-08-30
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PRIOR FILING DATE: 1999-08-30
NUMBER OF SEQ ID NOS: 7615
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              TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAACACAGCATGAGAACACAG
                                                          CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
                                                                                                                       GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG
                                                                                                                                                                                   CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
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CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/151,422
PRIOR FILING DATE: 1999-08-30
NUMBER OF SEQ ID NOS: 3855
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 3346
LENGTH: 1925
TYPE: DNA
ORGANISM: Homo Sapiens
US-09-652-917-3346
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GENERAL INFORMATION:
APPLICANT: Shyjan, Andrew W.
APPLICANT: Holtzman, Douglas A.
APPLICANT: Distefano, Peter
ITILE OF INVENTION: NOVEL NUCLEIC ACID
ITILE OF INVENTION: THEREFOR
ITILE REFERENCE: 1600.1170-001
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                                                                                                                      CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
                                                                                                                                                                  GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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RESULT 29
US-09-652-128-9375/c
US-09-652-128-9375/c
Sequence 9375, Application US/09652128
GENERAL INFORMATION:
APPLICANT: Shyjan, Andrew W.
APPLICANT: Shyjan, Andrew W.
APPLICANT: Holtzman, Douglas A.
FIITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
FITTLE OF INVENTION: HERREFOR
FILLE REFERENCE: 1600.1171-001
CURRENT APPLICATION NUMBER: US/09/652,128
CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/151,133
PRIOR APPLICATION NUMBER: 60/151,133
PRIOR FILING DATE: 1999-08-30
NUMBER OF SEO ID NOS: 10265
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LENGTH: 1925
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Best Local
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Pred. No. 4.4e-110;
0; Mismatches 1;
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APPLICANT: Curtis, Rory A.J.

APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: Distefano, Peter
ITILE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
ITILE OF INVENTION: THEREFOR
FILE REFERENCE: 1600.2020-001
CURRENT APPLICATION NUMBER: US/09/699,997
CURRENT FILING DATE: 2000-10-30
PRIOR APPLICATION NUMBER: 60/162,359
PRIOR APPLICATION NUMBER: 10-29
PRIOR FILING DATE: 1999-10-29
VOMBER OF SEQ ID NOS: 12714
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11085
LENGTH: 1925
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; ORGANISM: Homo sapiens
US-09-699-997-11085
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                                                                                                                                  CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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                     TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480
                                                                                      GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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 TAAATAAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-710-281-4458
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US-09-710-281-4458/c
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SOFTWARE: Fast
; SEQ ID NO 4458
FENGTH: 1925
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Best Local
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Hunter, John J.

APPLICANT: Stydan, Andrew W.

APPLICANT: Stbdal, Hilde

TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES

TITLE OF INVENTION: THEREFOR

FILE REFERENCE: 1600.2036-001

CURRENT APPLICATION NUMBER: US/09/710,281

CURRENT APPLICATION NUMBER: US/09/710,281

CURRENT FILING DATE: 2000-11-10

PRIOR APPLICATION NUMBER: 60/164,254

PRIOR FILING DATE: 1999-11-09

NUMBER OF SEQ ID NOS: 5803

SOFTWARE: FRASSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                       336 CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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                               GGTACCAATTTITCAGTGTTGGACTTGACAGCITCTTTTGCCACAAGCAAGAGAGAATTT
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TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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Pred. No. 4.4e-110;
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PRIOR FILING DATE: 2000-01-20
NUMBER OF SEQ ID NOS: 156
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 38
LENGTH: 1953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; NAME/KEY: SITE
; LOCATION: (1362)
; OTHER INFORMATION: n equals a,t,g, or
PCT-US00-09066-38
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PCT-US00-09066-38/c
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APPLICANT: Human Genome Sciences, In
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Best Local :
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CURRENT FILING DATE: 2000-04-06
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PRIOR FILING DATE: 1999-04-09
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                                             TGGGCAGACACGAGCAGGAGGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG
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Pred. No. 4.3e-110;
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US-09-726-805-1608/c
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SEQ ID NO 1608
LENGTH: 1925
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APPLICANT: Holtzman, Douglas A.
TITLE OF INVENTION: MOVEL NUCLEIC ACID MOLECULES AND USES
TITLE OF INVENTION: THEREFOR
FILE REFERENCE: 1600.2017-001
CURRENT APPLICATION NUMBER: US/09/726,805
CURRENT FILING DATE: 2000-11-30
PRIOR APPLICATION NUMBER: 60/168,140
PRIOR FILING DATE: 1999-11-30
NUMBER OF SEQ ID NOS: 2158
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TGGGCAGACACGAGCAGGAGGGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG
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 Mismatches

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Db 84 ACAGTG 79

RESULT 35

US-09-950-033-1914/c
Sequence 1914, Application US/09950083
GENERAL INFORMATION
APPLICANT: ROSEN, et. al
TITLE OF INVENTION: Human Secreted Proteins
FILE REFERENCE: RS805
CURRENT APPLICATION NUMBER: US/09/950,083
CURRENT APPLICATION NUMBER: 09/278,650
PRIOR PHILING DATE: 2001-03-12
PRIOR APPLICATION NUMBER: PCT/USO0/0604
PRIOR FILING DATE: 2001-04-12
PRIOR FILING DATE: 2001-09-12
PRIOR PHILING DATE: 2001-09-12
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2000-09-26 NUMBER: PCT/U	BER: P	NUMBER: PC	NUMBER: PCT/US00, 2000-06-01	2000-06-01	2000-06-01	2000-06-01	NUMBER: P	NUMBER: P	NUMBER 2000-	2000-06-01	2000-06-01	2000-06-01 NUMBER: PCT/USO	NUMBER: PC	NUMBER: PC	NUMBER: PC	NUMBER: PC 2000-04-0	2000-04-0	2000-04-06	2000-04-06	2000-04-06 NUMBER: PCT/USO	NUMBER: PC	NUMBER: P	NUMBER: PC	NUMBER: PC 2000-04-0	2000-04-0	0-03-23	0-03-23	2000-03-23	00-03-23 BER: PCT/USO	2000-03-2 NUMBER: PC	NUMBER: PC	NUMBER: P	NUMBER: PC	MBER: PC	000-03-23	000-03-23	R: PC	MBER: PC	BER: PC	3-2	MBER: PCT/US
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AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG

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Matches Query Match Best Local  $\vdash$ AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 605; Similarity Conservative 88.8%; 99.8%; 0; Score 555; DB 36; Pred. No. 4.3e-110; Mismatches Indels 0; Gaps 60 625 0,

Length 1953;

US-10-105-299-2414

LOCATION: (1362)..(1362) OTHER INFORMATION: n equals a,t,g,

or

FEATURE:
NAME/KEY: misc\_feature TYPE: DNA
ORGANISM: Homo sapiens

Query Match Best Local Matches

Similarity

99.8%;

Score 555; DB 40; Pred. No. 4.3e-110; Mismatches

Length 1953; Indels

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Gaps

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Conservative

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NUMBER OF SEQ ID NOS: 15197
NUMBER OF SEQ ID NOS: 15197
Prior Application removed SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2414
LENGTH: 1953

PRIOR PRIOR PRIOR FILING DATE: 1999-12-03 APPLICATION NUMBER: US 60/126,051 FILING DATE: 1999-03-23 APPLICATION NUMBER: US (FILING DATE: 1999-03-19 APPLICATION APPLICATION NUMBER: US FILING DATE: 1999-12-03 APPLICATION NUMBER: US 60/168,666 NUMBER: US 60/168,664 60/169,906 60/125,359

FILING DATE: 1999-03-12
APPLICATION NUMBER: US 60/138,597
FILING DATE: 1999-06-11

PRIOR PRIOR FILING DATE: 1999-03-12
APPLICATION NUMBER: US 60/138,574
FILING DATE: 1999-06-11
APPLICATION NUMBER: US 60/168,667 FILING DATE: 1999-12-03
APPLICATION NUMBER: US 60/124,142

PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR FILING DATE: 1999-06-11
APPLICATION NUMBER: US 60/168,662
FILING DATE: 1999-12-03
APPLICATION NUMBER: US 60/124,144

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PRIOR PRIOR PRIOR PRIOR PRIOR OR FILING DATE: 1999-12-03
OR APPLICATION NUMBER: US 60/124,095
OR APPLICATION NUMBER: US 60/138,598
OR FILING DATE: 1999-05-11
OR APPLICATION NUMBER: US 60/168,665
OR FILING DATE: 1999-12-03
OR FILING DATE: 1999-12-03
OR FILING DATE: 1999-12-03
OR APPLICATION NUMBER: US 60/125,360
OR APPLICATION NUMBER: US 60/125,360
OR APPLICATION NUMBER: US 60/138,626

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361 384

US-10-105-299-2414/c

Sequence 2414, Application US/10105299 GENERAL INFORMATION:

APPLICANT: Rosen, et.

a1

CURRENT APPLICATION NUMBER: US/10/105,299 TITLE OF INVENTION: Human Secreted Proteins FILE REFERENCE: PS950

2002-03-26

See File Wrapper or

CURRENT FILING DATE:

PRIOR PRIOR PRIOR FILING DATE: 1999-03-12
APPLICATION NUMBER: US 60/168,663

PRIOR PRIOR PRIOR PRIOR PRIOR APPLICATION NUMBER: US 60/168,622 FILING DATE: 1999-12-03 APPLICATION NUMBER: US 60/124,143 FILING DATE: 1999-12-03
APPLICATION NUMBER: US 60/124,096 FILING DATE: 1999-03-12

PRIOR PRIOR PRIOR APPLICATION NUMBER: US 60/168,661 FILING DATE: 1999-12-03 FILING DATE: 1999-03-12

PRIOR PRIOR FILING DATE: 1999-12-03
APPLICATION NUMBER: US 60/124,099

PRIOR PRIOR PRIOR PRIOR FILING DATE: 1999-03-12
APPLICATION NUMBER: US 60/168,654 FILING DATE: 1999-11-23
APPLICATION NUMBER: US 60/124,145

PRIOR PRIOR PRIOR PRIOR APPLICATION NUMBER: US 60/124,093 FILING DATE: 1999-03-12 APPLICATION NUMBER: US 60/166,989

PRIOR FILING DATE: 1999-03-12
APPLICATION NUMBER: US 60/167,061
FILING DATE: 1999-11-23 APPLICATION NUMBER: US 60/124,146 FILING DATE: APPLICATION NUMBER: 2000-09-26 2001-04-27 PCT/US01/13318

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CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAAGGGGAG

CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG

CAGCGACAGGCGGCAGCACCAGCACCTGCACGAACCCCGCCGAAACCTGCTGCGAGGACAC CAGCGACAGGCGCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC

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GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC

444 241 504 181 564 121

PRIOR PRIOR PRIOR FILING DATE: 2000-09-26
APPLICATION NUMBER: PCT/US00/26323
FILING DATE: 2000-09-26 APPLICATION NUMBER: PCT/US00/26337

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APPLICANT: HUMAN GENOME Sciences, Inc.
APPLICANT: Human Genome Sciences, Inc.
FITLE OF INVENTION: 49 Human Secreted Proteins
FILE REFERENCE: PS559PCT
CURRENT APPLICATION NUMBER: PCT/US00/14973
CURRENT FILING DATE: 2000-06-01
EARLIER APPLICATION NUMBER: 60/138,630
EARLIER FILING DATE: 1999-06-11
NUMBER OF SEQ ID NOS: 160
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 23
SEQ ID NO 3
LENGTH: 1956
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: SITE
NAME/KEY: SITE
LOCATION: (1362)
OTHER INFORMATION: n equals a,t,g, or c
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                                                                                                                                                                                                               Query Match
Best Local Similarity
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                                             GCCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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                                                                                                                                                                                          Score 555; DB 1;
Pred. No. 4.3e-110;
0; Mismatches 1;
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PRIOR APPLICATION NUMBER: 60/278,650
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: 09/833,245
PRIOR APPLICATION NUMBER: PCT/US01/11988
PRIOR FILING DATE: 2001-04-12
PRIOR PILING DATE: 2001-04-12
PRIOR APPLICATION NUMBER: PCT/US00/06043
PRIOR FILING DATE: 2000-03-09
PRIOR APPLICATION NUMBER: PCT/US00/06043
PRIOR APPLICATION NUMBER: PCT/US00/06058
PRIOR APPLICATION NUMBER: PCT/US00/06044
PRIOR APPLICATION NUMBER: PCT/US00/06044
PRIOR APPLICATION NUMBER: PCT/US00/06044
PRIOR APPLICATION NUMBER: PCT/US00/06059
PRIOR APPLICATION NUMBER: PCT/US00/06042
PRIOR APPLICATION NUMBER: PCT/US00/06042
PRIOR APPLICATION NUMBER: PCT/US00/06013
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PRIOR APPLICATION NUMBER: PCT/US00/06013
PRIOR APPLICATION NUMBER: PCT/US00/06049
PRIOR PILING DATE: 2000-03-09
PRIOR APPLICATION NUMBER: PCT/US00/06049
PRIOR PILING DATE: 2000-03-09
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PRIOR FILING DATE: 2000-03-09
PRIOR FILING DATE: 2000-03-09
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US-09-950-083-1915/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/950,083
CURRENT FILING DATE: 2001-09-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Human Secreted Proteins FILE REFERENCE: PS805
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NUMB 199	-06-	1999-03-19	MBER: U	MBER: US	 ⊃ ::	9-03-12	1999-12-03	MBER: US	NUMBER: US 60/124,143	1999-12-03	999-03-12	MBER: US	NUMBER: US	NUMBER: US 60/124,099 : 1999-03-12	1999-12-03	1999-03-12	MBER: US	NUMBER: US	: 1999-03-12	1999-11-23	8EM	NUMBER: US	NUMBER: PC	-60	2000-09-26	00-09-26 BER: PCT/	BER: PC	BER: PC	NUMBER: PC	06-01	2000-06-01	M C	NUMBER: PC	MBER: P	8	2000-06-01	BM	MBER: P	: 2000-06-01	2000-06-01	MBER: PC	MBER: PC	: 2000-04-06	2000-04-06	MBER: P

NUMBER: PCT/US00/08980

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RESULT 39
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; Sequence 2415, Application US/10105299
; GENERAL INFORMATION:
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OR FILING DATE: 1999-03-12
OR APPLICATION NUMBER: US 60/138,574
OR FILING DATE: 1999-06-11
OR APPLICATION NUMBER: US 60/168,667
OR FILING DATE: 1999-12-03
OR APPLICATION NUMBER: US 60/124,142
OR FILING DATE: 1999-03-12
OR APPLICATION NUMBER: US 60/138,597
OR APPLICATION NUMBER: US 60/168,666
OR FILING DATE: 1999-10-10
OR APPLICATION NUMBER: US 60/168,666
OR FILING DATE: 1999-12-03
OR FILING DATE: 1999-12-03
OR APPLICATION NUMBER: US 60/168,666
OR FILING DATE: 1999-12-03
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OR FILING DATE: 1999-03-19
OR APPLICATION NUMBER: US 60/168,664
OR FILING DATE: 1999-12-03
OR APPLICATION NUMBER: US 60/126,051
OR APPLICATION NUMBER: US 60/126,050
OR APPLICATION NUMBER: US 60/169,906
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Pred. No. 4.3e-110;
0; Mismatches 1; Indels 0
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LENGTH: 1956
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1362)...(1362)
; OTHER INFORMATION: n equals a.
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NUMBER OF SEQ ID NOS: 15197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Rosen, et. al TITLE OF INVENTION: Human Secreted Proteins FILE REFERENCE: PS950 CURRENT APPLICATION NUMBER: US/10/105,299 CURRENT FILING DATE: 2002-03-26
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ACAGTG
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                                    CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
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Pred. No. 4.3e-110;
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RESULT 40 PCT-US02-29964-16

Sequence 16, Application PC/TUS0229964
GENERAL INFORMATION:
APPLICANT: Tang, Y. Tom
APPLICANT: Ren, Feiyan

٠	· ADDITCANT: Zhang, Jie		
•••	APPLICANT: Zhao, Qing A.		
	<pre>; APPLICANT: Wang, Jian-kui : APPLICANT: Xue, Aidong J.</pre>		
	APPLICANT: Wang, Dunrui		
	; APPLICANT: Asundi, Vinod		
	; APPLICANT: Wehrman, Tom		
٠. ٠	; APPLICANT: Goodrich, Ryle W.		
	; APPLICANT: Weng, Gezni ; APPLICANT: Haley-Vicente, Dai	na	
	APPLICANT: Drmanac, Radoje T	io Poide	
•• •	; TITLE OF INVENTION: Polypept	es actus an	
	; FILE REFERENCE: BUYACIP PCT : CURRENT APPLICATION NUMBER: PC		
•• •	CURRENT FILING DATE: 2002-09	9	
·· ··	<pre>; PRIOR APPLICATION NUMBER: US       PRIOR FILING DATE: 2001-09-19</pre>		
•••	PRIOR APPLICATION NUMBER: US	09/488,725	
٠. ٠.	; PRIOR APPLICATION NUMBER: US	09/552,317	
••	FILING DATE: 2000-04-25		
	APPLICATION NUMBER: PCT FILING DATE: 2000-12-22	/0500/35017	
••	; PRIOR APPLICATION NUMBER: US	09/491,404	
	APPLICATION NUMBER: PCT	/US01/02623	
• •	: PRIOR FILING DATE: ZOUI-U1-ZO	09/496.914	
	; PRIOR FILING DATE: 2000-02-03	7560.8	
٠.	; PRIOR FILING DATE: 2000-04-27		
٠. ٠.	; PRIOR FILING DATE: 2001-02-05	501/038	
	; PRIOR APPLICATION NUMBER: US ; PRIOR FILING DATE: 2000-02-28	09/515,126	
	; Remaining Prior Application d. ; NUMBER OF SEO ID NOS: 992	ata removed - See File Wrapper or PALM.	
• • • •	So	6.0	
•• ••	; SEQ ID NO IO ; LENGTH: 1967		
	: ORGANISM: Homo saniens		
٠. ٠	FEATURE:		
g }	; NAME/KEY: CDS ; LOCATION: (338)(763)		
D.	CT-U		
	Query Match  Best Local Similarity 99.8%;  Matches 605; Conservative	Score 555; DB 1; Length 1967; Pred. No. 4.3e-110; 0; Mismatches 1; Indels 0; Gap	ps
ρ	1 AGTTCTCCTTGCAGAGGACTGG	GCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG	60
₽	1349 AGTTCTCCTTGCAGAGGACTGG	CGCCGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG	1408
ρ	61 GGCGCTGTCGGTGGAGTGC	CATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG	120
Дb	1409 GGCGCTGTCGGTGGAGTGC		1468
Qy	121 CAGCGACAGGCGGCAGCACAGC	CTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC	180
g	1469 CAGCGACAGGCGGCAGCAGA	CCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC	1528
δ	181 CGTGTACAGGAGCGGGTTGATG	CGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG	240
Вр	1529 CGTGTACAGGAGCGGGTTGATG	ACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG	1588
Ş	241 GAGGATCATGTACGCCCGGAAG	AGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC	300
밁	1589 GAGGATCATGTACGCCCGGAAG	TAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC	1648

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GENERAL INFORMATION:
APPLICANT: MOITIS, MacDonald
APPLICANT: Lal, preeti
APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide Polymorp
FILE REFERENCE: GX-0006 P
CURRENT APPLICATION NUMBER: US/60/172,373
CURRENT APPLICATION NUMBER: US/60/172,373
CURRENT FILING DATE: 1999-12-16
NUMBER OF SEQ ID NOS: 25,772
SEQ ID NO 15734
LENGTH: 716
                                                                                                                                                                                                           Matches
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Best Local :
                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: 227484.3c
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AGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATG
                           CGGCAGCACGAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGG
                                                        CGGCAGCACAGCACCTGCACGAACACCCCGCCGAAACTGCTGCGAGGACACCGTGTACAGG 190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACC
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                                                                                                                                                                                                            595;
                                                                                                                                                                                                                     h 87.2%;
Similarity 99.8%;
                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                   Score 545; DB o1,
Pred. No. 8e-108;
                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                               DB 61; Length 716;
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APPLICANT: Xu, Jiangchun
APPLICANT: Xtolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
APPLICANT: Molesh, David Alan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.48401PC
CURRENT APPLICATION UNMBER: PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEO ID NOS: 215
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LENGTH: 1619;
TYPE: DNA;
ORGANISM: Homo sapiens
PCT-US01-45395-205
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PCT-US01-45395-205
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                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
                                 1112
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      240 GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAG
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                                                                                                                                                                   61 GGCGCTGTCGGTGGTGGAGTGCGCGATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                                                                                                                                    Match 77.8%; Local Similarity 99.8%;
                                                                                                                                                                                                                                         1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
                                                                                                                       CAGCGACAGGCGGCAGCACAGCACC-TGCACGAACACCCGGCGAAACTGCTGCGAGGACA 179
                                                                                           GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 1111
                                                                                                                                                                                                              AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 1051
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Pred. No. 2.7e-95;
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SEQ ID NO 205
LENGTH: 1619
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C4
CURRENT FILING DATE: 200-11-14
NUMBER: US/09/713,550
CURRENT FILING DATE: 200-11-14
NUMBER OF SEQ ID NOS: 205
                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1352 CTGGGCAGACACGAGCAGGAGGGAGAGACAGAGAAAAGAAAAACACAGCATGAGAACACA 1411
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              300 CCATGATCCTCCGAATCTGGTTGGGCATTCCAGCATACGGCCAATGTCACAACAATCAGCC
                                                                                                                                                                                              121 CAGCGACAGGCGGCAGCACAGCACC-TGCACGAACACCCGGCGGAAACTGCTGCGAGGACA 179
                                                                                                                                                                                                                                      Match 77.8%;
Local Similarity 99.8%;
                                                                                                                                                                                                                                                                                                                              1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
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                                                                       GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGGTTTGGCCGCAG 299
                                                                                                                                                                                                                                                                                                  CCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC
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                                                                                                                                                                                                                                                                                                                                                       Score 486; DB 28; Pred. No. 2.7e-95; 0; Mismatches 0;
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1351
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ğ	1232	GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGGGTTTGGGTTTGGCCGCAG 1291
8 8	300 1292	CCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC 359
8 8	360 1352	CTGGGCAGACACGAGCAGGAGGAGAGACACAGAAAAAAAA
8 8	420 1412	GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA 479 
₽ ₹	480 1472	TGGTACCAATTTTTCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAG
8 5	540 1532	TAACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAG
8 5	600 1592	GACAGTG 506         acagtg 1598
RESU Se GE A A	47640000	15-211 (Applic) FORMATION: COTIXA C COTIXA C Xu, Jia Stolk, Algate, Fling,
CH S S S C C F H H	12"4 PAZZA	LE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY LE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER E REFERENCE: 20121.48401PC RENT APPLICATION NUMBER: PCT/US01/45395 RENT FILING DATE: 2001-11-13 BER OF SEQ ID NOS: 215 TWARE: PastSEQ for Windows Version 4.0 ID NO 211 NOTH: 1619 PE: DNA GANISM: Homo sapiens G1-45395-211
Qu Be	ery Ma st Loc tches	tch 77.8%; Score 486; DB 1; Length 1619; al Similarity 99.8%; Pred. No. 2.7e-95; 606; Conservative 0; Mismatches 0; Indels 1; Gaps
용 성	1 992	AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
유 첫	61 1052	GCCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
β 5.	121 1112	CAGCGACAGGCGGCAGCACAGCACC-TGCACGAACACCCCGCCGAAACTGCTGCGAGGACA 179 
8 5	180 1172	CCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAAACGTCTCCGAGAAGGGGA 239 
β 5⁄ <sub>2</sub>	240 1232	GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCCTTGGGTTTGGCCGCAG 299

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Search completed: November 8, 2002, 01:59:57 Job time: 1449.81 secs

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/cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*
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US-10-264-237-1380
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US-09-721-456-3
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US-09-721-456-3
US-09-721-75-208
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US-10-127-850A-533
US-10-128-689A-533
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US-10-131-833A-533
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Sequence 208, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 533, App Sequence 534, App Sequence 534, App Sequence 534, App Sequence 534, App Sequence 534, App Sequence 534, App Sequence 534
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US-10-240-453-117	US-10-264-237-1221	US-10-264-237-447	US-10-264-237-1069	US-10-264-237-662	US-08-250-795A-19	US-10-137-873A-533	US-10-137-872A-533	US-10-131-836A-533	US-10-131-821A-533	US-10-128-693A-533	US-10-128-690A-533	US-10-128-686A-533	US-10-128-685A-533	US-10-128-684A-533	US-10-127-849A-533	US-10-127-843A-533	US-10-127-838B-533	US-10-127-825A-533
Sequence 117, App		447	1069		19,	533,	•	533,	533,	533,	533,	533,	533,	Sequence 533, App	•	•		Sequence 533, App

ALIGNMENTS

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RESULT 1

US-10-230-437-15

Sequence 15, Application US/10230437

GENERAL INFORMATION:

APPLICANT: Baker, Kevin P.
APPLICANT: Besnoyers, Luc
APPLICANT: Desnoyers, Luc
APPLICANT: Goddard, Audrey
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APPLICANT: Goddard, Colon
APPLICANT: Godder, 1097-09-17
APPLICANT: Godder, 1097-09-17
APRIOR APPLICATION NUMBER: GO/062287
APRIOR APPLICANTION NUMBER: GO/06287
APRIOR FILING DATE: 1997-10-18
APRIOR APPLICANTION NUMBER: GO/078910
APRIOR APPLICANTION NUMBER: GO/078910
APRIOR FILING DATE: 1998-03-20
APRIOR FILING DATE: 1998-03-20
APRIOR FILING DATE: 1998-03-26
APRIOR APPLICATION NUMBER: GO/079284
APRIOR APPLICATION NUMBER: GO/079284
APRIOR APPLICATION NUMBER: GO/079284
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APR
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62.9%;

Score 393;

DB 6;

Length 1524;

Matches

1132

1192

61

121

Best Local Similarity

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; NAME/KEY: misc_feature; LOCATION: (1297)..(1297); OTHER INFORMATION: n equals a,t,g, or US-10-264-237-1044
Sequence 1380, Application US/10264237
GENERAL INFORMATION:
APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA131P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1044, Application US/10264237 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
SOFTWARE: PATCHIN Ver. 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Nucleic Acids, Proteins, FILE REFERENCE: PA1311 CURRENT APPLICATION NUMBER: US/10/264,237 CURRENT FILING DATE: 2002-10-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1252 CAGCGACAGGCGGCAGCACCAGCACCTGCACGAAACACCCGCCGAAACTGCTGCGAGGACAC 1311
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100.0%; Pr
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Pred. No.
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                         and Antibodies
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                                          ; FEATURE:
NAME/KEY: misc_feature
LOCATION: (1998)..(1998)
OTHER INFORMATION: n equals
US-10-264-237-464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 4
US-10-264-237-464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver.
SEQ ID NO 1380
LENGTH: 2049
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 464, Application US/10264237 GENERAL INFORMATION:
                                                                                                                                                                                                          SOFTWARE: Pa
SEQ ID NO 464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
      Query Match
                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
PRIOR FILING DATE: 2000-05-19
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NUMBER OF SEQ ID NOS: 2876
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PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR APPLICATION NUMBER: US 60/205,515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PA131P1
                                                                                                                                                                                                                                                  PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Birse et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (12)..(12)
OTHER INFORMATION: n equals
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LOCATION: (1819)..(1819)
OTHER INFORMATION: n equals
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (1987)..(1987)
OTHER INFORMATION: n equals a,t,g,
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                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                    TYPE: DNA
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                                                                                                                                                                                     ENGTH:
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nes 22; Conserv
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Pred. No.
          Score 21;
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        Length 2062;
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US-10-264-237-1044

APPLICANT: Birse et al.

RESULT 3 US-10-264-237-1380/c

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Matches Best Query Match

Local

Similarity

SEQ ID NO 1044

ENGTH:

1329

FEATURE:

ORGANISM: Homo sapiens

TYPE: DNA

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RESULT 5
US-09-721-456-3
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                                                                                    TELEFAX: (415) 576-0
INFORMATION FOR SEQ ID NO: /no
SEQUENCE CHARACTERISTICS:
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                                                                                                                                          TELEPHONE: (415) 576-0200
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Human Telomerase Catalytic Subunit NUMBER OF SEQUENCES: 727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGNAAAAAAAAAAAAAAAAA 625
                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                              NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
                                                                                                                                                                                                                                                        APPLICATION NUMBER: WO PCT/US97/17885 FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/974,549A FILING DATE: 19-NOV-1997 APPLICATION NUMBER: US 08/724,643 FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/721,456 FILING DATE: 22-Nov-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: WO PCT/US97/17618 FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: California
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                                                                LENGTH: 2176 base pairs
                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 08/915,503 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 08/912,951 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/911,312 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 08/854,050 FILING DATE: 09-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/851,843 FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/846,017 FILING DATE: 25-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
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Harley, Calvin B.
Andrews, William H.
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Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ingner, Joachim
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                                                                                                                          576-0300
                                                                                                      /note= "clone 712562"
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; OTHER INFORMATION: Genbank Accession No. AA631399
US-10-240-425-208
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                                    SOFTWARE: Patent.pm
SEQ ID NO 21763
LENGTH: 163
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SEQ ID NO 208
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Best Local Similarity
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Best Local
                                                                                                                                                                                          APPLICANT: Giordano, J.Y.

TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.

FILE REFERENCE: 59.US2.REG
                                                                                              PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/240,425
CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: PCT/US01/09847
PRIOR FILING DATE: 2001-03-28
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APPLICANT: Scherf, Uwe
APPLICANT: Vockley, Joseph G.
TITLE OF INVENTION: Gene Expression Profiles in
FILE REFERENCE: 44921-5026
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TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
ORGANISM: Homo sapiens
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LOCATION: 1..2176
SEQUENCE DESCRIPTION: SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alvarez,
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Lord, Reginald V.
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Pred. No.
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Pred. No.
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0.29;
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0.91;
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; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: 135
; OTHER INFORMATION: n-a,
US-09-513-999C-21763
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                                                                                        Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 550 SEQ ID NO 533
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Best Local Similarity
Matches 19; Conserv
                                                                                                                                            PRIOR EILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR EILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C139
CURRENT APPLICATION NUMBER: US/10/131,813A
CURRENT FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: 60/049911
PRIOR PELLING DATE: 1997-06-18
PRIOR PELLING DATE: 1997-06-18
PRIOR PELLING DATE: 1997-08-26
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-08-26
                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
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PRIOR TILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
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LOCATION: 134
OTHER_INFORMATION: w=a
                                  TYPE: DNA
ORGANISM: Homo Sapien
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NAME/KEY: unsure
                       FEATURE:
                                                                       LENGTH: 496
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Godowski, Paul J.
Gurney, Austin L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Watanabe, Colin K
Wood, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Smith, Victoria
Stewart, Timothy A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tumas, Daniel
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Pred. No. 2.8;
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                                                                                                                             See File Wrapper or PALM
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                                                     ; FEATURE:
; NAME/KEY: unsure
; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-131-819A-533
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US-10-131-819A-533
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                                                                                                                                                                                     SEQ ID NO 533
LENGTH: 496
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 Best Local Similarity
                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
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PRIOR APPLICATION NUMBER: 60/059115
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PRIOR FILING DATE: 1997-06-18
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PRIOR APPLICATION NUMBER: 60/059352
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                                                                                                                                                TYPE: DNA ORGANISM: Homo Sapien
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                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/059117 FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                     FILING DATE: 1997-09-
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/059122
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Filvaroff, Ellen
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 3.0%;
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L00.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                         60/059263
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 Score 19; pred. No.
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 DB 6;
2.7;
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                   Length 496;
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RESULT 10
US-10-131-823A-533
Commence 533, App
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                                                 Query Match
Best Local Similarity
Watches 19; Conserve
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                                                                                                                                    ; OTHER INFORMATION: unknown base US-10-131-823A-533
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LENGTH: 496
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APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maur
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PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C143
                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/131,823A CURRENT FILING DATE: 2002-04-24 PRIOR APPLICATION NUMBER: 60/049911 PRIOR FILING DATE: 1997-06-18 PRIOR APPLICATION NUMBER: 60/056974 PRIOR FILING DATE: 1997-08-26 PRIOR APPLICATION NUMBER: 60/059113 PRIOR APPLICATION NUMBER: 60/059113
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                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1997-09-19
                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/059588
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                                                                                                                                                                  NAME/KEY: unsure
LOCATION: 396
                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                   ORGANISM: Homo Sapien
                                                                                                                                                                                                                                       TYPE: DNA
396 NAAAAAAAAAAAAAAAA 414
                               607 NAAAAAAAAAAAAAAAAA 625
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Watanabe, Colin K
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Godowski, Paul J.
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                                                                 Conservative
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                                                                               3.0%; Score 19;
100.0%; Pred. No.
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RESULT 12
US-10-131-826A-533
; Sequence 533, Application US/10131826A
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
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; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-131-824A-533
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US-10-131-824A-533
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PRIOR FILING DATE: 1997-06-18
PRIOR PPLICATION NUMBER: 60/056974
PRIOR PPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR PPLICATION NUMBER: 60/059113
PRIOR PPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059117
PRIOR PILING DATE: 1997-09-17
PRIOR PILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059127
PRIOR APPLICATION NUMBER: 60/059127
PRIOR APPLICATION NUMBER: 60/059127
PRIOR PILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
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Best Local Similarity 100.0%; I
Matches 19; Conservative 0;
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LENGTH: 496
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
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ORGANISM: Homo Sapien
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                                                                                                                                                                           607 NAAAAAAAAAAAAAAAAAA 625
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Pred. No.
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Beresini, Maureen

Desnoyers,Luc DeForge,Laura

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CURRENT APPLICATION NUMBER: US/10/131,826A
CURRENT FILING DATE: 2007-04-24
PRIOR APPLICATION UMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR FILING DATE: 1997-06-674
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION UMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          data removed - See File Wrapper or PALM
                                                                                                                                                                                                                                                                                                                                          Score 19; pred. No.
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US-10-125-926A-533; Sequence 533, App
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C138
CURRENT APPLICATION NUMBER: US/10/131,829A
CURRENT FILING DATE: 2002-04-27
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APPLICANT:
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PRIOR
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PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
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ORGANISM: Homo Sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: unsure LOCATION: 396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                  607 NAAAAAAAAAAAAAAAAA 625
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Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
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                                   Sherwood, Steven
                                                       Gurney, Austin L.
                                                                       Goddard, Audrey
Godowski, Paul J.
                                                                                                            Gerritsen, Mary E.
                                                                                                                                Gao, Wei-Qiang
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Filvaroff, Ellen
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100.0%; Pred. No.
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR APPLICATION NUMBER: 60/059184
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
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APPLICATION NUMBER:

APPLICATION NUMBER: 60/059117 FILING DATE: 1997-09-17

APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: PPLICANT:

Zhang, Zemin

Tumas, Daniel Watanabe, Colin K Wood, William Stewart, Timothy A. Godowski, Paul J. Gurney, Austin L. Sherwood, Steven

Goddard, Audrey Gerritsen, Mary E. Gao, Wei-Qiang Filvaroff, Ellen

Smith, Victoria

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; OTHER INFORMATION: unknown base US-10-131-826A-533

NAME/KEY: unsure LOCATION: 396

FEATURE:

ORGANISM: Homo Sapien

SEQ ID NO 533 LENGTH: 496

NUMBER OF SEQ ID NOS: 550

PRIOR APPLICATION NUMBER: 60/059588 PRIOR FILING DATE: 1997-09-19

TYPE: DNA

Query Match Best Local

Local

h 3.0%; Similarity 100.0%; 19; Conservative (

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Matches

19;

RESULT 13 US-10-131-829A-533

Sequence 533, Application US/10131829A GENERAL INFORMATION:

APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Mauree
APPLICANT: Desnoyers, Luc
APPLICANT: Desnoyers, Luc
APPLICANT: filvaroff, Eller
APPLICANT: Gao, Wei-Oilang
APPLICANT: Gao, Wei-Oilang
APPLICANT: Gerritsen, Mary

Desnoyers, Luc Filvaroff, Ellen Beresini, Maureen

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RESULT 15
US-10-127-829A-533
Sequence 533, Application US/10127829A
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                                       APPLICANT:
                                                                         APPLICANT:
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PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
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ORGANISM: Homo Sapien
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nes 19; Conserv
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FILING DATE: 1997-09-18
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FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059122
FILING DATE: 1997-09-17
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FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059115
FILING DATE: 1997-09-17
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     OF INVENTION:
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                     T: Zhang, Zemin
INVENTION: SECR
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                                                      Watanabe, Colin K
Wood, William
                                                                                      Stewart, Timothy A. Tumas, Daniel
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Godowski, Paul J.
Gurney, Austin L.
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SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ACIDS ENCODING THE SAME
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100.0%; Pred. No.
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US-10-127-831A-533
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
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Best Local
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                 CURRENT APPLICATION NUMBER: US/10/127,831A CURRENT FILING DATE: 2002-10-15 PRIOR APPLICATION NUMBER: 60/049911 PRIOR FILING DATE: 1997-66-18
                                                                                     TITLE OF INVENTIÓN: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C107
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PRIOR FILING DATE: 1997-08-26
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RIOR APPLICATION NUMBER: 60/056974
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FILING DATE: 1997-09-18
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FILING DATE: 1997-09-17
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Sherwood, Steven
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 495
TYPE: DNA
ORGANISM: Homo Sapien
                                                                 CURRENT APPLICATION NUMBER: US/10/127,835A CURRENT FILING DATE: 2002-10-15 PRIOR APPLICATION NUMBER: 60/049911 PRIOR FILING DATE: 1997-06-18 PRIOR APPLICATION NUMBER: 60/056974 PRIOR FILING DATE: 1997-08-26 PRIOR APPLICATION NUMBER: 60/059113 PRIOR APPLICATION NUMBER: 60/059113
                                                                                                                                                                                                                              TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C102
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100.0%; Pred. No.
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US-10-127-837A-533
Cacuence 533, Application US/10127837A
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CURRENT APPLICATION NUMBER: US/10/127,837A
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/049911
PRIOR TILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059120
PRIOR APPLICATION NUMBER: 60/059120
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PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
Remaining Prior Application data remonumber of SEQ ID NOS: 550
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LENGTH: 496
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NAME/KEY: unsure
LOCATION: 396
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APPLICATION NUMBER: 60/059122
FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059184
FILING DATE: 1997-09-17
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19; Conserv
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Wood,William
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US-10-127-831A-533

NAME/KEY: unsure LOCATION: 396 OTHER INFORMATION: unknown base

FEATURE

PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19

PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/ PRIOR FILING DATE: 1997-09-17 PRIOR APPLICATION NUMBER: 60/

60/059352

PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-17

60/059117 60/059115 60/059113

FILING DATE: 1997-0: APPLICATION NUMBER: FILING DATE: 1997-0 APPLICATION NUMBER:

1997-09-17 1997-08-26

PRIOR APPLICATION NUMBER: 60/059122 PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER:

60/059184 60/059263

Matches Query Match

Local Similarity es 19; Conserva

Conservative

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RESULT 17 US-10-127-835A-533 ; Sequence 533, Application US/10127835A

GENERAL INFORMATION:

Baker, Kevin P.

APPLICANT:

APPLICANT: APPLICANT:

APPLICANT:

Gao, Wei-Qiang Filvaroff, Ellen Desnoyers, Luc DeForge, Laura Beresini, Maureen

APPLICANT:

APPLICANT:

Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.

APPLICANT: APPLICANT: APPLICANT:

APPLICANT: APPLICANT: APPLICANT:

Watanabe, Colin K Wood, William Zhang, Zemin

Tumas, Daniel Stewart, Timothy A Smith, Victoria Sherwood, Steven

PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17

NUMBER: 60/059117

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PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
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CURRENT FILING DATE: 2002-10-15
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PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-19
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LOCATION: 396
OTHER INFORMATION: unknown base
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ORGANISM: Homo Sapien
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                         APPLICATION NUMBER: 60/059588 FILING DATE: 1997-09-19
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                                                             FILING DATE: 1997-09-19
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FILING DATE: 1997-09-17
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                                                                                                                 APPLICATION NUMBER: 60/059263
       Prior Application
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100.0%; Pred. No.
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data removed - See File Wrapper or PALM.
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RESULT 20
US-10-127-850A-533
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SEQ ID NO 533
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SEQ ID NO 533
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C110
CURRENT APPLICATION NUMBER: US/10/127,850A
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR APPLICATION NUMBER: 60/049911
PRIOR APPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-08-26
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
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                                                                         NUMBER OF SEQ ID NOS: 550
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PRIOR FILING DATE: 1997-09-19
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TYPE: DNA
ORGANISM: Homo 9
TYPE: DNA
ORGANISM: Homo Sapien
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                                         LENGTH: 496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: unsure
LOCATION: 396
OTHER INFORMATION: unknown base
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Local Similarity 100.0%;
                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Baker, Kevin P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumas, Daniel
Watanabe, Colin K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wood, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stewart, Timothy A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Smith, Victoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sherwood, Steven
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                                                                                                             1997-09-19
                                                                                                                             60/059588
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9-17
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                                                                                        data removed
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; Pred. No.
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    See File Wrapper or PALM.

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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C86
CURRENT APPLICATION NUMBER: US/10/127,901A
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/05974
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/05912
PRIOR APPLICATION NUMBER: 60/05912
PRIOR APPLICATION NUMBER: 60/05912
PRIOR APPLICATION NUMBER: 60/05912
PRIOR FILING DATE: 1997-09-17
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; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-127-850A-533
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                                         ; FEATURE:
; NAME/KEY: unsure
; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-127-901A-533
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US-10-127-901A-533
                                                                                                                                                                          PRIOR FILING DATE: 1997-09-19
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
TYPE: DNA
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Best Local
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Query Match
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es 19; Conserv
                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 60/059352
FILING DATE: 1997-09-19
APPLICATION NUMBER: 60/059588
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Godowski, Paul J.
Gurney, Austin L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Smith, Victoria
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Filvaroff, Ellen
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100.0%; Pred. No. 2.
Live 0; Mismatches
  3.0%;
    Score 19;
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    DB 6;
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Length 496;
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US-10-128-689A-533
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CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR ETLING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
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Matches 19; Conserv
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LENGTH: 496
                                    Matches
                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1997-09-19
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PRIOR APPLICATION NUMBER:
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                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo Sapien
FEATURE:
                                                                                                                                              NAME/KEY: unsure LOCATION: 396
                                                                                                                               OTHER INFORMATION: unknown base
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607 NAAAAAAAAAAAAAAAAA 625
                                                       Local Similarity
                                      19;
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Wood, William
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                                      Conservative
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                                100.08; +-
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100.0%; Pred. No.
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                                        Mismatches
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                                                                           DB 6;
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                                                                         Length 496;
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                                        Indels
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APPLICANT:

Baker, Kevin P.

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RESULT 24
US-10-131-833A-533
: Sequence 533, Application US/10131833A
: GENERAL INFORMATION:
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NAME/KEY: unsure
; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-131-830A-533
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US-10-131-830A-533
                                                                                                                                                                                                                                                                                                                  Remaining Prior Application data removed - See File Wrapper NUMBER OF SEQ ID NOS: 550 SEQ ID NO 533 LENGTH: 496 TYPE: DNA
                                                                                                                                                                           Query Match
Best Local (
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GENERAL INFOR
                                                                                                                                                            Matches
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
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PRIOR FILING DATE: 1997-09-19
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PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
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                                                                                                            607 NAAAAAAAAAAAAAAAAA 625
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FILING DATE: 1997-09-19
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Godowski, Paul J.
Gurney, Austin L.
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1997-09-17
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RESULT 25
US-10-131-837A-533
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CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
                                                                               Sequence 533, Application US/10131837A GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
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NUMBER OF SEQ ID
SEQ ID NO 533
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Best Local :
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                                   APPLICANT:
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PRIOR FILING DATE: 1997-09-19
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PRIOR FILING DATE: 1997-09-19
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                   APPLICANT:
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nes 19; Conserv
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Gao, Wei-Qiang
              Desnoyers, Luc
Filvaroff, Ellen
                                                               Beresini, Maureen
                                                 DeForge, Laura
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Wood, William
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Filvaroff, Ellen
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D NOS: 550
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APPLICANT:

Godowski, Paul J. Gurney, Austin L. Gerritsen, Mary E.

Goddard, Audrey

APPLICANT:

APPLICANT:

APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT:

APPLICANT:

Zhang, Zemin Wood, William Watanabe, Colin K Tumas, Daniel Stewart, Timothy A Smith, Victoria Sherwood, Steven

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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REPERENCE: P33301C131
CURRENT APPLICATION NUMBER: US/10/131,837A
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/05974
PRIOR APPLICATION NUMBER: 60/059113
PRIOR ETLING DATE: 1997-08-26
PRIOR ETLING DATE: 1997-08-26
PRIOR FILING DATE: 1997-09-17
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US-10-127-825A-533
; Sequence 533, Applic; GENERAL INFORMATION:
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LENGTH: 496
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Best Local Similarity
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TITLE
                                        APPLICANT:
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  OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
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                                    Watanabe, Colin K
Wood, William
                                                                                             Smith, Victoria
Stewart, Timothy A.
                                                                                                                                       Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
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                                                                              Tumas,Daniel
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100.0%; Pred. No.
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; FEATURE: NAME/KEY: unsure ; LOCATION: 396 ; OTHER INFORMATION: unknown base US-10-131-837A-533

SEQ ID NO 533 LENGTH: 496

TYPE: DNA

ORGANISM: Homo Sapien

PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19

PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263

PRIOR APPLICATION NUMBER: 60/059122 PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/059184

PRIOR APPLICATION NUMBER: 60/059115 PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/059588 PRIOR FILING DATE: 1997-09-19

RESULT 26 US-10-125-930A-533

Sequence 533, Applic GENERAL INFORMATION

Application US/10125930A

APPLICANT: APPLICANT:

> DeForge, Laura Beresini, Maureen

APPLICANT:

APPLICANT: Baker, Kevin P.

APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT:

Goddard, Audrey Godowski, Paul J. Gurney, Austin L. Sherwood, Steven

Smith, Victoria

Gao, Wei-Qiang Gerritsen, Mary E. Desnoyers, Luc Filvaroff, Ellen 밁 õ

Best Query Match

Local 19;

Similarity 100.0%;

3.0%; Score 19; 00.0%; Pred. No 0;

DB 6;

Mismatches No.

0;

Conservative

Matches

APPLICANT:

Stewart, Timothy A.

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RESULT 28
US-10-127-838B-533
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; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-127-825A-533
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CURRENT APPLICATION NUMBER: US/10/127,838B
CURRENT FILING DATE: 2002-04-22
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                   APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C98
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PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
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PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
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CURRENT APPLICATION NUMBER: US/10/127,825A
CURRENT FILING DATE: 2002-04-22
PRIOR ADDITORNOON
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NUMBER OF SEQ ID NOS: 550
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PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
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ORGANISM: Homo Sapien
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Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
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Watanabe, Colin K
Wood, William
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
TYPE: DNA
ORGANISM: Homo Sapien
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                                                                 CURRENT APPLICATION NUMBER: US/10/127,843A
CURRENT FILING DATE: 2002-04-22
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
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APPLICANT:
                                  PRIOR APPLICATION NUMBER: 60/059113 PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                          TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C99
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PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
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PRIOR APPLICATION WASTER 1997-09-17
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
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PRIOR FILING DATE: 1997-08-26
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NAME/KEY: unsure
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APPLICATION NUMBER: 60/059115 FILING DATE: 1997-09-17
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19; Conserv
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Wood, William
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Gurney, Austin L.
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Filvaroff, Ellen
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100.0%; Pred. No.
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US-10-127-843A-533
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
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US-10-127-849A-533
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PRIOR FILING DATE: 1997-09-19
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550

SEQ ID NO 533
LENGTH: 496
TYPEP - TYPEP.
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GENERAL INFORMATION
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PRIOR FILING DATE: 1997-09-19
                                  PRIOR APPLICATION NUMBER: 60/59115
PRIOR FILING DATE: 1997-09-17
PRIOR PPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                           TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3390R10103

CURRENT APPLICATION NUMBER: U5/10/127,849A

CURRENT FILING DATE: 2007-04-23

PRIOR APPLICATION NUMBER: 60/049911

PRIOR FILING DATE: 1997-06-18

PRIOR FILING DATE: 1997-06-18
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LOCATION: 396
OTHER INFORMATION: unknown base
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ORGANISM: Homo Sapien
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APPLICATION NUMBER: 60/
FILING DATE: 1997-09-17
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Wood, William
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Godowski, Paul J.
Gurney, Austin L.
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100.0%; Pred. No.
tive 0; Mismatc
                60/059184
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2.7;
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; OTHER INFORMATION: unknown base
US-10-127-849A-533
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US-10-128-684A-533
                                                            PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
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PRIOR FILING DATE: 1997-06-18
PRIOR PPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR PILING DATE: 1997-09-17
PRIOR PILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059117
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
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NAME/KEY: unsure
NAME/KEY: unsure
NAME/KEY: 396
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PRIOR EILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR EILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C118
CURRENT APPLICATION NUMBER: US/10/128,684A
CURRENT FILING DATE: 2002-04-23
CURRENT FILING DATE: 2002-04-23
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                                           APPLICATION NUMBER: 60/
APPLICATION NUMBER: 60/
                              FILING DATE: 1997-09-19
APPLICATION NUMBER: 60/059588
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Query Match Best Local (

FEATURE:

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RESULT 32
US-10-128-685A-533
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                                     Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 550 SEQ ID NOS: 533 LENGTH: 496
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENCTH: 496
TYPE: DNA
ORGANISM: Homo Sapien
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PRIOR APPLICATION NUMBER: 60/0
PRIOR FILING DATE: 1997-09-19
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P333071C1116
TO TO TO THE SAME
FILE REFERENCE: P333071C1116
TO TO TO TO THE SAME
FILE REFERENCE: P333071C1116
TO TO TO TO THE SAME
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                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/059113
PRIOR TILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
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TYPE: DNA
ORGANISM: Homo Sapien
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LOCATION: 396
OTHER INFORMATION: unknown base
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Godowski, Paul J.
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100.0%; Pred. No.
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US-10-128-686A-533
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; NAME/KEY: unsure
; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-128-685A-533
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                                                                                                                                                          Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 550 SEQ ID NO 533
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Best Local Similarity 100.0%;
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PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
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PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C119
CURRENT APPLICATION NUMBER: US/10/128,686A
CURRENT FILING DATE: 2002-04-23
CURRENT FILING DATE: 2002-04-23
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
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            OTHER INFORMATION: unknown base
                                NAME/KEY: unsure LOCATION: 396
                                                                                           TYPE: DNA
ORGANISM: Homo Sapien
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                                                                                                                                        LENGTH: 496
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FILING DATE: 1997-09-17
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Wood, William
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Godowski, Paul J.
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Filvaroff, Ellen
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100.0%; Pred. No.
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2.7;
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396 NAAAAAAAAAAAAAAAAA 414

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RESULT 34
US-10-128-690A-533
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
                                                                                           Query Match
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C122
CURRENT APPLICATION NUMBER: US/10/128,690A
CURRENT FILING DATE: 2002-04-23
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR FILING DATE: 1997-06-18
PRIOR FILING DATE: 1997-06-18
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/0
PRIOR FILING DATE: 1997-09-17
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PRIOR APPLICATION NUMBER: 60/059352
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PRIOR FILING DATE: 1997-09-17
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                                                                                                                                               NAME/KEY: unsure
LOCATION: 396
OTHER INFORMATION: unknown base
                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo Sapien
                                                                                                                                                                                                       FEATURE:
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                   607 NAAAAAAAAAAAAAAAAA 625
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                                                    Local Similarity es 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 1997-09-19
Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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0; Mismatches
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Pred. No. 2.7;
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RESULT 35
US-10-128-693A-533
RESULT 36
US-10-131-821A-533
; Sequence 533, Application US/10131821A
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
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PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
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                                                                                                                                                                                                                                                       LOCATION: 396
OTHER INFORMATION: unknown base
                                                                                                                                                                                                                                                                                          NAME/KEY: unsure
                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo Sapien
                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                LENGTH: 496
                                                                                            396
                                                                                                               607 NAAAAAAAAAAAAAAAAA 625
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L00.0%; Pred. No.
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INFORMATION:

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RESULT 37
US-10-131-836A-533
; Sequence 533, Application US/10131836A
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
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PRIOR APPLICATION NUMBER: 60/0499
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/0569
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/0591
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
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Best Local
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LENGTH: 496
APPLICANT:
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LOCATION: 396
OTHER INFORMATION: unknown base
                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo Sapien
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                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/059122 FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 1997-09-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60 FILING DATE: 1997-09-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Baker, Kevin P.
                           Beresini, Maureen
DeForge, Laura
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Wood, William
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Stewart, Timothy A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Smith, Victoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sherwood, Steven
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Godowski, Paul J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application data removed - See File Wrapper or PALM ) NOS: 550\,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER: 60/059117
1997-09-17
                                                                                                                                                                                                                                                               3.0%; Score 19; 100.0%; Pred. No.
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                                                                                                                                                                                                                                              0;
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Sequence 533, Applic GENERAL INFORMATION:

Application US/10137872A

APPLICANT: Baker, Kevin P. APPLICANT: Beresini, Maur

APPLICANT: APPLICANT: APPLICANT:

APPLICANT:

Goddard, Audrey Godowski, Paul J. Gurney, Austin L.

Sherwood, Steven

APPLICANT:

APPLICANT:

Desnoyers, Luc Filvaroff, Ellen

Beresini, Maureen DeForge, Laura

Gao, Wei-Qiang Gerritsen, Mary E. APPLICANT:

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RESULT 38
US-10-137-872A-533
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PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 533
LENGTH: 496
                                                                                                                                            Matches
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PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 1997-0
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                                                                                                                                                                                                                                                                                                                                                                            Remaining Prior A
NUMBER OF SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER:
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                                                                                                                                                                                                                          NAME/KEY: unsure
LOCATION: 396
OTHER INFORMATION: unknown base
                                                                                                                                                                                                                                                                                     ORGANISM: Homo Sapien FEATURE:
                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                        607 NAAAAAAAAAAAAAAAAA 625
                                                                      396 NAAAAAAAAAAAAAAAAA 414
                                                                                                                                            Local Similarity
nes 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059117
FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059122
FILING DATE: 1997-09-17
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Wood, William
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Godowski, Paul J.
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                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                             Application data removed - See File Wrapper or PALM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1997-09-17
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                                                                                                                                                           3.0%; Score 19;
100.0%; Pred. No.
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                                                                                                                                            0;
                                                                                                                                            Mismatches
                                                                                                                                                           DB 6;
2.7;
                                                                                                                                          0
                                                                                                                                                                          Length 496;
                                                                                                                                          Indels
                                                                                                                                          0;
                                                                                                                                          Gaps
                                                                                                                                         0
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Tumas, Daniel Smith, Victoria

Wood, William Watanabe, Colin K Stewart, Timothy A.

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PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR PELICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 533, Application US/10137873A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/137,872A CURRENT FILING DATE: 2002-05-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.0%; Score 19;
100.0%; Pred. No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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Best Local S
Matches 19
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PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19, Application US/08250795A GENERAL INFORMATION:
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LENGTH: 496
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PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 1997-09-19 Remaining Prior Application d NUMBER OF SEQ ID NOS: 550
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PRIOR FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Homo Sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: unsure
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     396 NAAAAAAAAAAAAAAA 414
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Vincent, Matthew P.
REGISTRATION NUMBER: 36,70
                                                                                                                                        COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII (text)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: ROPES & GRAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Immunosuppressant Target Proteins NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Berlin, Vivian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . Similarity
19; Conserv
                                                                                         APPLICATION NUMBER: US/08/250,795A FILING DATE: 27-May-1994
                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 02110
                                                                                                                                                                                                                                                                                                                                       STATE: MA
                                                                                                                                                                                                                                                                                                                                                           STREET: 1 International Place CITY: Boston
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100.0%; Pred. No.
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2.7;
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Query Match Best Local S Matches 19

Local Similarity

Conservative

100.0%; F tive 0;

US-10-137-873A-533

GENERAL INFORMATION:

Baker, Kevin P.

APPLICANT:

Tumas,Daniel Watanabe,Colin K Wood,William

APPLICANT APPLICANT APPLICANT: APPLICANT: APPLICANT:

APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT:

> Gerritsen, Mary E. Gao, Wei-Qiang Desnoyers, Luc Filvaroff, Ellen DeForge, Laura Beresini, Maureen

Goddard, Audrey Godowski, Paul J. Gurney, Austin L. Sherwood, Steven

Smith, Victoria

Stewart, Timothy A.

APPLICANT:

; OTHER INFORMATION: unknown base US-10-137-872A-533

NAME/KEY: unsure LOCATION: 396

PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17

PRIOR FILING DATE: 1997-09-19

LENGTH: 49 TYPE: DNA

ORGANISM: Homo Sapien

FEATURE:

PRIOR FILING DATE: 1997-09-18 PRIOR APPLICATION NUMBER: 60/059263

APPLICATION NUMBER: 60/059184 FILING DATE: 1997-09-17

APPLICATION NUMBER: 60/059352 FILING DATE: 1997-09-19

APPLICATION NUMBER: 60/059588

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TOPOLOGY: linear;
; MOLECULE TYPE: cDNA
; SEQUENCE DESCRIPTION: SEQ ID NO: 19:
US-08-250-795A-19
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Sequence 662, Application US/10264237

GENERAL INFORMATION:

APPLICANT: Birse et al.

TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA131P1

CURRENT APPLICATION NUMBER: US/10/264,237

CURRENT ETLING DATE: 2002-10-04

PRIOR FILING DATE: 2001-05-18
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NUMBER OF SEQ ID NOS: 2876
SOFTWARE: Patentin Ver. 3.1
SEQ ID NO 662
LENGTH: 692
TYPE: DNA
                                                                  Matches
                                                                            Query Match
Best Local Similarity
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SEQUENCE CHARACTERISTICS:
LENGTH: 531 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                         NAME/KEY: misc_feature
LOCATION: (692)..(692)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (651)..(651)
OTHER INFORMATION: n equals
                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
LOCATION: (23)..(23)
OTHER INFORMATION: n equals a,t,g, or c
                                                                                                                                                                                                           LOCATION: (684)...(684)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (684)..(684)
                                                                                                                                                                                                                                                                                                                                                 LOCATION: (32)..(32)
OTHER INFORMATION: n equals
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                             607 NAAAAAAAAAAAAAAAA 625
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651 NAAAAAAAAAAAAAAAAA 669
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 951-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         . Similarity
19; Conserv
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                                                           3.0%; So ilarity 100.0%; I Conservative 0;
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ilarity 100.0%;
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Pred. No.
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2.7;
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Sequence 1069, Application US/10264237

GENERAL INFORMATION:
APPLICANT: Birse et al.
ITITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA131P1
CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
SOFTWARE: Patentin Ver. 3.1
SEQ ID NO 1069
LENGTH: 1330
RESULT 44
US-10-264-237-1221
; Sequence 1221, Appli
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                    ; FEATURE:

NAME/KEY: misc_feature

; LOCATION: (2046)..(2046)

; OTHER INFORMATION: n equals a,t,g, or

US-10-264-237-447
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US-10-264-237-447
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SEQ ID NO 447
                                                                                                                                                                                  Matches
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Best Local S
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CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA131P1
                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 2876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (1265)...(1265)
OTHER INFORMATION: n equals a,t,g, or
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ORGANISM: Homo sapiens
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19; Conserv
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Pred. No.
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APPLICANT: Birse et al.

Application US/10264237

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PRIOR FILING DATE: 2001-04-06
PRIOR FILING DATE: 2001-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR APPLICATION NUMBER: DE 10019173.8
PRIOR FILING DATE: 2000-04-07
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 350
SEQ ID NO 117
LENGTH: 15832
                                    , LUCATION: (5221, 5244, US-10-240-453-117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-240-453-117/c
US-10-240-453-117/c

Sequence 117, Application US/10240453

GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
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  Query Match
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Best Local .
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SOFTWARE: PATENTIN VER. 3.1
SEQ ID NO 1221
LENGTH: 3196
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Diagnosis of Diseases Associated with DNA
TITLE OF INVENTION: Transcription
TITLE OF INVENTION: by Means of Assessing the Methylation Status of Genes Associated
TITLE OF INVENTION: with DNA Transcription
FILE REFERENCE: 5013.1009
CURRENT PRICTON NUMBER: US/10/240,453
CURRENT FILING DATE: 2002-10-02
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CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (3170)..(3170)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (668)..(668)
OTHER INFORMATION: n equals a,t,g,
                                                                      NAME/KEY: unsure LOCATION: (5221,
                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                             OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                        FEATURE:
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LOCATION: (791, 2400,
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                                                                                                                                         3281, 4994, 5001..5002, 5006,
                                                                      6508, 12917, 12968, 12987..12988, 13055;
    3.0%;
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Pred. No. 2.5;
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  DB 6;
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BQ006545 UI-H-EII-
AII336826 wp69h10.x
BF439382 nab63g06.
BF594242 7n10d03.x
AII990500 ws40d07.x
BM669397 UI-E-DW1-
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AW128849 xd46f12.x	Q689771 AGENCO	363261 qy56f02.	66797 tn23q08.x	3057775 7£75a10	73739 zp04h05.r	F726644 by09q04	084249 zn17f09	075710 zm89d04.	970361 op52q	65640 aq55d12.	3162 tf32c12.x	AW191974 x178a02	72281 ap23c03.x	125134 6017	126050 60176288	AW087372 xh19g02	42092 wa38h03 x	054680 7169f08	BF054837 7171h01.	65236 w172h08 x	350014 ht07g12	630 to03b07	91683 gv93f04	613995 no86c08	1769 ov326	M667957 UT-E-DW	150931 gh53c04 y	85990 60	686 wl83d07	75598 vh24e07	W149665 xf40h05	36858 gy86g11 v	W338938 ha68	F726459 bv06h03	F939693 nac80h12	58216 7g19g07	90950 hg51e12	7680 ACENCON

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## ALIGNMENTS

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FEATURES SOUTCE	JOURNAL COMMENT	TITLE	REFERENCE	SOURCE ORGANISM	ACCESSION VERSION KEYWORDS	RESULT 1 BQ006545/c LOCUS DEFINITION
Email: cgapbs-r@mail.nih.gov Tissue Procurement: Dr. Jose Mercuende cDNA Library preparation: Dr. M. Bento Soares, University of Iowa cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa DNA Sequencing by: Dr. M. Bento Soares, University of Iowa Clone Distribution: Clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Seq primer: M13 FORWARD POLYA-Yes. Location/Qualifiers 1702	Tould bene index Unpublished (1997) Contact: Robert Strausberg, Ph.D.	NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP),	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 702)	human. Homo sapiens	IMAGE:5846228 3', mRNA sequence. BQ006545 BQ006545.1 GI:19731445 EST.	BO006545  702 bp mRNA linear EST 26-MAR-2002 UI-H-EIl-aza-n-21-0-UI.sl NCI_CGAP_EIl Homo sapiens cDNA clone

/clone="IMAGE:5846228" /clone\_lib="NCI\_CGAP\_EI1" /tissue\_type="Chondrosarcoma"

/organism="Homo sapiens" /db\_xref="taxon:9606"

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445 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
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                                                                                                                                                                                                                                        TGGGCAGACACGAGCAGGAGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note-"Organ: Left Pelvis: Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not NCI_CGAP_EII is a normalized cDMA library containing the following tissue(s): Chondrosarcoma. The library was
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/lab_host="DH10B (Life Technologies)"
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1 (bases 1 to 678)

NCI/NIMOS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project
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                                                                                                                                                                                                                                                                                                                               This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                       Tumor Gene Index
Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria;
1 (bases 1 to 696)
                                                                                                                                                                                                                                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
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Mammalia; Eutheria; Primates;
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/note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from five normalized libraries were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 5 libraries. The pools consisted of the following libraries and cloneIDs: Soares NDSHF pool 1: 145032-147335, 147720-148103, 148872-149255, 15002 - 150407, 151176-152327 Soares ND2HF8-9W pool 1: 758280-760583, 772104-774407 Soares NDHPA pool 1: 304776-306311, 320136-322823, 326280-326663 Soares NbHOT
                                                                                                                                                                                                                                                                                                                  quality sequence stop: 459
                                                                                                                                                                                                             /clone="IMAGE:3272627"
/clone_lib="Soares_NSF_F8_9W_OT_PA_P_S1"
/lab_host="DH10B"
                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
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1 (bases 1 to 537)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index Unpublished (1998)

Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                GPR39. ;, mRNA sequence.
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7n10d03.x1 NCI_CGAP_Brn23 Homo sapiens ci
similar to SW:GP39_HUMAN 043194 PUTATIVE
Tissue Procurement: David N. Louis, M.D.,
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Location/Qualifiers
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
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/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                     27 GGGACGCGAAGAGCAACGGGCGCTGCACAAAGCGGGGCGCTGTCGGTGGAGTGCGCAT 86
                                                                                                                                                                                                                               ACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCA
                                    ACCTCGTCCAGTCGTGCTTGGGTTTGGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCA
                                                                                         AGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGG
                                                                                                                        AGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGG
                                                                                                                                                           GCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCG
                                                                                                                                                                             GCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCG
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Score 529; Pred. No. 0; Mismatches

DB 9;

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RESULT 5 AI990500/c

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> 117 490 177 430 237 370 297 310 357 250 417 190 477 130

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ws40d07.x1 NCI_CGAP_GC6 Homo sapiens cDNA clone IMAGE:2499661 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 1212 Std Error: 0.00
Seq primer: -40UP from Gibco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop:
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R. Emmert-Buck, M.D., Ph.D.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (bases 1 to 616)
/note="Vector: pT/T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; plasmid DNA from the normalized ilbrary NCI_CGAP_GC4 was prepared, and so circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified CDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1469064-1470983, and 1475592-1476743). Subtraction by Bento Soares and M. Fatima Bonaldo. " a Subtraction by Bento Soares and M. Fatima Bonaldo."
                                                                                                                                                                                                                                                                                                                                                    /clone="INAGE:2499661"
/clone_lib="NCI_CGAP_GC6"
/tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
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FEATURES
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                                                                                                                                                                                                                                                                                                                                            Email: msoares@blue.weeg.uiowa.edu
Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
cDNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
Clone Distribution: Researchers may obtain clones from Research
Genetics (www.resgen.com).
Seq primer: M13 Forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               contact: Soares, MB
Program for Rat Gene Discovery and Mapping
University of Iowa
451 Eckstein Maddan.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 627)
Bonaldo,M.F., Lennon,G. and Soares,M.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BM669397 627 bp mRNA linear EST 27 UI-E-DW1-ahd-c-11-0-UI.sl UI-E-DW1 Homo sapiens cDNA clone UI-E-DW1-ahd-c-11-0-UI 3', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                   POLYA-Yes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EST
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319 335 9565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Res. 6 (9), 791-806 (1996)
/lab_host="DHIOB (Life Technologies) (T1 phage resistant)"
/note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a
modified polylinker; Site_1: EcoR I; Site_2: Not I;
UI-E-DW1 is a normalized cDNA library containing the
following tissue(s): lens. The library was constructed
according to Bonaldo, Lennon and Soares, Genome Research,
6:791-806, 1996. First strand cDNA synthesis was primed
with an oligo-dT primer containing a Not I site. Double
stranded cDNA was ligated to an EcoR I adaptor, digested
with Not I, and cloned directionally into pT7T3-pac
vector. The oligonucleotide used to prime the synthesis of
                                                                                                                                                                                            /tissue_type="lens"
/dev_stage="adult"
                                                                                                                                                                                                                             /clone="UI-E-DW1-ahd-c-11-0-UI"
/clone_lib="UI-E-DW1"
                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
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TGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG
                                                                                   AGCTTCTTTTGCCACAAGCAAGAGAGAATTTAACACTGTTTCAAACCCGGGGGAGTTGGC
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                      AACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTA 576
                                                                                                                                                                                                                                                                                                                                                                                                     CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCCAATGTCACCAACAATCAGCCC 360
                                                                                                                                                                                                                                    TAAATGAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTTGCTG 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCCTGCTG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 124
                                                                                                                                                                                     TGGGCAGACACGAGCAGGAGGAGAGAGAGAAAAGAAAAACACAGCATGAGAACACAG 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 240
                                                                                           CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
AACACTGTTTCAAACCCGGGGGGAGTTGGCTGTGTTA 640
                                                                                                                                                                                                                                                                                   CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM12722 row: h column: 07
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Tissue Procurement: Invitrogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 quality sequence stop: 641.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="organ: brain; Vector: pCMV-SPORT6; Site_1: EcoRV (destroyed); Site_2: Not1; RNA source male hippocampus, age 27. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.4 kb, insert size range 0.9-4 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 012."

160 c 197 g 134 t 17 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5727798"
/clone_lib="NIH_MGC_124"
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/lab_host="DH10B"
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99.8%;
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Pred. No. 0;
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TITLE
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Best Local
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                                                                                                                                                                                                                                                                        568 CAACGGGCGCTGCACAAAGCGGGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCG 509
                                                                                                                                                                                                                                                                                               40 CAACGGGCGCTGCACAAAGCGGGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCG 99
ANACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTC
                                                                                                                                     CCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAA 219
                                                                                                                                                                                CTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGCCAGCACCAGCACCTGCACGAACACCCG
                                                                                          CCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAA
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hg51e12.x1 NCI_CGAP_GC6 Homo sapiens cDNA clone IMAGE:2949166 3'
similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Possible reversed clone: similarity Seq primer: -40UP from Gibco High quality sequence stop: 457.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LIANL at: image.llnl.gov/image/html/iresources.shtml
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bonaldo, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota;
Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tissue Procurement: Christopher A. Moskaluk, M.D., R. Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: pT773D-Pac (Pharmacia) with a modified /note="Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Plasmid DNA from the normalized library NCI_CGAP_GC4 was prepared, and so circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1469064-147098), and 1475592-1476743). Subtraction by Bento Soares and M. Fatima Bonaldo. "
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/clone="IMAGE:2949166"
/clone_lib="NCI_CGAP_GC6"
/tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. 0;
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                                                                                                                                                                                                                                                                                                                               (CGAP/BTGAP), Tumor Gene Index Unpublished (1998)
Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis,
                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostor Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 551)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
                                                                                                                                                                                                                                              Bonaldo, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7g19g07.x1 NCI_CGAP_Brn23
similar to SW:GP39_HUMAN (
                                                                                                                                                                                     High quality sequence stop: 490 Location/Qualifiers
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                                                                                                                                                                                                                           info@image.llnl.gov
                                                                                                                                                                                                                                                                                                       cDNA Library Preparation: M. Bento
                                                                                                                                                                                                               primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens
                                                                                                                                                                                                                                       through the I.M.A.G.E. Consortium/LLNL, send email
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;, mRNA sequence.
/clone="IMAGE:3306972"
/clone=11b="NCI_CGAP_Brn23"
/tlssue_type="911oblastoma (pooled)"
/lab_host="DH10B"
                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       551 bp mRNA linear EST 29-SEP-2000
3 Homo sapiens cDNA clone IMAGE:3306972 3'
043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
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                                                                                                                                                                                                                                                                                                                                M.D., Myrna
                                                                                                                                                                                                                                                                                                       Soares,
                                                                                                                                                                                                                                                                                                         ph.D.,
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae
1 (bases 1 to 540)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of N
ational Cancer Institute / National Institute of N
Disorders and Stroke, Brain Tumor Genome Anatomy Pr
(CGAP/BTGAP), Tumor Gene Index
Unpublished (1998)
Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis,
                                                                                                                                                                                                                                                                                                                                                   BF939693 540 bp mRNA linear EST 22-JAN-2001 nac80bl2.xl NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:3440591 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
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cDNA Library Arrayed by: Greg Lennon, Ph.D.

cDNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
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onaldo, Ph.D.
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Location/Qualifiers
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/db_xref="taxon:9606"
/clone="IMAGE:3440591"
/clone_lib="NOI_CGAP_Brn23"
/tlssue_rype="glioblastoma (pooled)"
/lab_host="DH10B"
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99.8%;
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AGGAGGGAGAGACAGAGAAAAAAAAACACAGGATGAGAAACACAGTAAATGAATAAAACC 435 	CTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGC 375 	CCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAAT 315	GTTGATGACCGAGCTGAGGTAGAAAAAGGTCTCCGAGAAGGGGAGGAGGAGGATCATGTACGC 255 	GCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGG 195	/ Match 75.4%; Score 471; DB 12; Length 530; Local Similarity 100.0%; Pred. No. 0; nes 471; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	/tissue_type="Lens" /dev_stage="Adult" /lab_host="EMDHIDB" /lab_rost="EMDHIDB" /note="Organ: Eye; Vector: pCMVSPORT6; Two human lenses from different adults (both approximately 40 years old) together yielded 20ug of total RNA and 150ng mRNA for cDNA library synthesis. A directionally cloned cDNA library in the pcMVSPORT6 vector was constructed at Life Technologies, essentially following the protocols of the SuperScript plasmid System full details of which are contained in the manufacturer's Instruction manual (http://www.lifetech.com/). First strand synthesis was carried out using a Not I primer-adapter [5'-pcACTACTTCTACATCGGGAGGGGCGCCCC(T)15-3']. Not I/blunt end inserts were cloned into the Not I/EcoR V sites in the vector. EST analysis was performed on the unamplified library at the NIH Intramural Sequencing Center (NISC)."	National Eye Institute 6/331, NIH, Bethesda, Tel: 301 402 3452 Fax: 301 496 0078 Email: graeme@helix.n Plate: 06 row: h co seq primer: M13RP1 re Location/Qua 1530 /organism="H /db_xref="ta /clone="by06 /clone=lib=" By"	<pre>imates; Catarrhini; Hominidae; Homo. ,S., Behal,A. and Smith,D. and bioinformatics for ocular genomics s. Sci. 41, (2000) In press tructure and Function</pre>	uman Lens cDNA (Un clone by06h03 5', 3I:12042370

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REFERENCE
AUTHORS
TITLE
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                                                                                       CACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAG
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                                                                                                                                                          CAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCAGCACAGCGCAGCACAGCACCTGCACGAA 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCGGGGGAGTTGGCTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG
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                                                                    CACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAG
                                                                                                                                         CAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCACCAGCACCTGCACGAA 468
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                                                                                                                                                                                                                512;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq
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1 (bases 1 to 527)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: ggapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AW338938 527 bp mRNA linear EST 31-JAN-2000 ha68h04.x1 NCI_CGAP_Pan1 Homo sapiens cDNA clone IMAGE:2878903 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
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                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Life Technologies catalog #: 11548-013
DNA Sequencing by: Washington University Genome Sequencing
                                                                                                                                                                                                                                                                                                      103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     quality sequence stop: 420
                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                   /note="Organ: pancreas; Vector: pCMV-SPORT6; Site_1: SalI;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.72 kb. Life Technologies catalog #:
                                                                                                                                                                                                                                                                                                                      11548-013"
                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="NCI_CGAP_Pan1"
/tissue_type="adenocarcinoma"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="IMAGE: 2878903"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R. Emmert-Buck, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                                                                                                                                                                                                                                                                    quality sequence stop: 444.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                   through the I.M.A.G.E. Consortium/LLNL
/note="Vector: PT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Plasmid DNA from the normalized library NCI_CGAP_GC4 was prepared, and so circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1469064-1470983, and 1475592-1476743).
                                                                                                                                                                               /clone_lib="NCI_CGAP_GC6"
/tissue_type="pooled germ
/lab_host="DH108"
                                                                                                                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2009444"
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97

Query Match Best Local S Matches 509

Similarity

73.48;

Conservative

BASE COUNT ORIGIN

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TTTGCCACAAGCAAGAGAGATTTAACACTGTTTCAAACCCGGGGAGTTGGCTGTGTTA
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 Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Burt Feuerstein, M.D., Mark Israel, M.D.
cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing C
Clone distribution: NCI-CGAP clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 690)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index (1998)

Opublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                     AW149665
                                                                                                                                                                                                                                                                                                                                                                                                                                        AW149665

690 bp mRNA linear EST 03-NOV-1999 xf40h05.xl NCI_CGAP_Brn50 Homo sapiens cDNA clone IMAGE:2620569 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                       AW149665.1 GI:6197561
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Pred. No. 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_"IMAGE:2620569"
/clone_lib="NCI_CGAP_Brn50"
/tissue_type="medulloblastoma"
/lab_host="DH10B (phage resistant)"
/lab_host="DH10B (phage resistant)"
/lab_host="DH10B (phage resistant)"
/note="Organ: brain; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from medulloblastoma tumor tissue, and was then primed with a Not I - ollgo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and to Eco RI adaptors (Pharmacia); digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. This library is normalized. Library constructed by Bento Soares and M. Fatima Bonaldo. "
42 a 196 c 160 g 191 t 1 others
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/db_xref="taxon:9606"
500 bp mRNA linear EST 13-OCT-1999
NCI_CGAP_Kidl3 Homo sapiens cDNA clone IMAGE:2577252 3'
SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
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              GCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGC
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                                                                               AGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAA 478
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Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: Christa Prange, The I.M.A.G.E. Consortium DNA Sequencing by: Washington University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 500)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Organ: kidney; Vector: pCMV-SPORT6; Site_1: Sal
Site_2: NotI; Cloned unidirectionally. Primer: Oligo
Library constructed by Life Technologies."
a 139 c 114 g 148 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="NGI_CGAP_Kid13"
/tissue_type="2 pooled Wilms' tumors, one primary and one
metastatic to brain"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:9606"
/clone="IMAGE:2577252"
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                                          1 AGTTCTCCTTGCAGAGGACTGGCGCGCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
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DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llni.gov/bbrp/image/image.html
Insert Length: 1462 Std Error: 0.00
Seq primer: -40UP from Gibco
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1 (bases 1 to 591)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1998)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      High quality sequence stop: 444.
Location/Qualifiers
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                                                                                                              Similarity
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                                                                                      Conservative
                                                                                                                                                                                              T 3']; double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized, and was constructed by Bento Soares and M.Fatima Bonaldo."

179 c 147 g 155 t 1 others
                                                                                                                                                                                                                                                                                                                                  /clone="IMAGE:2431501"
/clone_lib="NCI_CGAP_Brn25"
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                                                                                                                                                                                                                                                                                                                                                                                                                           /tissue_type="anaplastic oligodendroglioma"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
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100.0%;
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Pred. No. 0;
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GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120

Best Local Similarity

Pred. No.

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RESULT 17
BE385990
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EuKaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia, Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 676)
1 (bases 1 to 676)
NIH-MGC http://mgc.ncl.nlh.gov/.
NATIONAL Institutes of Health, Mammalian Gene Collect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
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                                                                                ECORI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit
                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:3617696"
/clone_lib="NIH_MGC_20"
                                                   (Stratagene) and 172 c 199
                                                                                                                                                                                      /tlssue_type="melanotic melanoma"
/lab_host="DH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB); Site_1: XhoI; Site_2:
                                                                                                                                                                                                                                                                                                                                    ocation/Qualifiers
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Query Match

Score

425;

DB

10;

Length

source

1. .467

/organism="Homo sapiens"
/db\_xref="taxon:9606"

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AUTHORS
TITLE
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1 (bases 1 to 467)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
                                                                          CONA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                    Ph.
                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
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                                                                                                                                                                                                                                        (CGAP/BTGAP),
                                                                                                                                                    cDNA Library Preparation: M. Bento Soares, Ph.D.,
                                primer: -40ml3 fwd. ET from Amersham
               quality sequence stop: 425
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                                                                                                                                      ph.D.
ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                              467 bp mRNA linear EST 26-OCT-1998
.NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:1703814 3'
.SW:NTR1_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1 ;, mRNA
                                                                                                                                                                                                                                                                                                                                                                                             GI:3679400
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                                                                                                                                                                                  Myrna R.
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                                                                                         Sequencing Center information can be
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AUTHORS
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MEDLINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               167
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                                                                                                                                                                                                                                                                                                                                                                                                                          ATGAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCCAGGAAATGGT 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCAGACACGAGCAGGAGGAGAGACAGAGAAAAAAAAACACAGCATGAGAACACAGTAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAG 408
                                                                                                                                                                                                                              UI-E-DWO-agm-j-01-0-UI.sl UI-E-DWO Homo sapiens cDNA UI-E-DWO-agm-j-01-0-UI 3', mRNA sequence.
                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 538)
Bonaldo, M.F., Lennon, G. and Soares, M.B.
                                                                                                                                                                                                  EST
Contact: Soares, MB
Program for Rat Gene Discovery and Mapping
                                                                     discovery
                                                                                 Normalization and subtraction: two
                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                BM667957.1 GI:18975788
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                                                  Res. 6 (9), 791-806 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T 3']; double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized, and was constructed by Bento Soares and M.Fatima Bonaldo."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="IMAGE:1703814"
/clone_lib="NCI_CGAP_Brn23"
/tissue_type="glioblastoma (pooled)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  67.7%;
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Pred. No. 0;
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                                                                               approaches
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Best Local 9
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                                                   191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                         CCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAA 314
                                                                                                                                                                                                                                                                                                                                                                                                                                           CATAAAATATTTAGCCCCTCTGTTCTGTGCTGTCTGGCCAGGAAATGGTACCAATTTTTC 494
                                                                                                                                                                       AGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAGAGAAATTTAACACTGTTTCAAA
                                                                                                                                             TCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAG
                                                                                                                                                                                                                                                                                           TCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAG
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Tissue Procurement: Dr. Gregg Hageman
CDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iow
CDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
Clone Distribution: Researchers may obtain clones from Research
Genetics (www.resgen.com).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          University of Iowa
451 Eckstein Medical Research Building Iowa City,
Tel: 319 335 8250
Fax: 319 335 9565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  POLYA=Yes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab_host="DHIOB (Life Technologies) (T1 phage resistant)"
/note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a
modified poolylinker; Site_1: EcoR I; Site_2: Not I;
UI-E-DWO is a cDNA library containing the following
tissue(s): lens. The library was constructed according to
Bonaldo, Lennon and Soares, Genome Research, 6:791-806,
1996. First strand cDNA synthesis was primed with an
oligo-dT primer containing a Not I site. Double stranded
cDNA was ligated to an EcoR I adaptor, digested with Not
I, and cloned directionally into pT7T3-Pac vector. The
oligonucleotide used to prime the synthesis of
first-strand cDNA contains a library tag sequence that is
located between the Not I site and the (dT).8 tall. The
sequence tag for this library is CGATTAGCGA. This library
was created for the program, Gene Discovery in the Visual
System, supported by National Eye Institute (NEI).
TAG TISSUFF human lens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ø
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TAG_TISSUE-human lens
TAG_SEQ=CGATTAGCGA"
144 c 122 g
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/clone_lib="UI-E-DW0"
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/db_xref="taxon:9606"
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99.88;
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Pred. No. 0;
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           71
                                                                                                                                                                                         84
                                    CCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGT 263
                                                                                              CCTGCACGAACACCCGCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGA 203
                                                                                                                                                 CATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGAGCAGGCGGCAGCAGCA 143
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CCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGT
                                                                          CCTGCACGAACACCCGCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGA 418
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AIO18769 537 bp mRNA linear EST 27-AUG-1998 ov32e04.xl Soares_testis_NHT Homo sapiens cDNA clone IMAGE:1639038 3' similar to SW:NTR1_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 537)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mRNA sequence.
AI018769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                www-bio.llni.gov/bbrp/image/image.html
Insert Length: 1394 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 452.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                                              105
                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                          Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT773 vector. Library went through one round of normalization to Cot5, and was constructed by Bento Soares and M. Fatima Bonaldo. " 147 c 126 g 159 t
                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:1639038"
/clone_lib="Soares_testis_NHT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex="male"
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                                                                                                                                                                                                                                                66.4%;
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                                                                                                                                                                                                                                                  Score 415;
Pred. No.
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                                                                                                                                                                                                                                                                    DB 9;
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similar to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Tissue Procurement: L. Jeffrey Medeiros, M.D., Michael
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                        High quality sequence stop: 395.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                 www-bio.llnl.gov/bbrp/image/image.html
Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                   DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                      56
                                                /clone_lib="NCI_CGAP_AA1"
                                                                                                                                                                                                                                                  /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
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SW:NTR_RAT P20789
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sapiens cDNA clone IMAGE:1113710 3'
NEUROTENSIN RECEPTOR ; mRNA sequence.
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David B. Krizman,

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 452)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological
Disorders and Stroke, Brain Tumor Genome Anatomy Project
(CGAP/BTGAP), Tumor Gene Index
                                                                                                                                                                                   cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 1183 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           452 bp mRNA linear EST 17-MAR-1999 qy93f04.x1 NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2019583 3'similar to SW:NTR1_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1 ;, mRNA
                                                                                                                                                                                                                                                                                                         Bonaldo,
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                                                                                                                                             quality sequence stop: 451.
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                /clone="IMAGE: 2019583"
/clone_lib="NCI_CGAP_Brn25"
                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
  /tissue_type="anaplastic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GACCATTAAATGCTTTAGACAGTG 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CACAGCATGAGAACACAGTAAATAAAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGT 146
                                                                                                                                                                                                              sequence.
            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Emmert-Buck, M.D., Ph.D.
                                                               Tumor Gene Index
Unpublished (1997)
                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 396)
                                                                                                                                                                                                                                  AI499630 396 bp mRNA linear EST 12-MAY-1999 to03b07.xl NCI_CGAP_Ut2 Homo sapiens cDNA clone IMAGE:2177941 3' similar to SW:NTRI_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1 ;, mRNA
                                                 Contact: Robert Strausberg, Ph.D.
                                                                                         NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Ana
                                                                                                                                                          Homo sapiens
cDNA Library Preparation: Life Technologies, Inc
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                                                                                                                                                                                                   GI:4391612
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99.8%;
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Pred. No. 0;
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                                                                                        Genome Anatomy Project (CGAP),
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BE350014/c
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 534)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="NCI_CGAP_Ut2"
/tissue_type="moderately-differentiated endometrial adenocarcinoma, 3 pooled tumors"
/lab_host="DH10B"
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/db_xref="taxon:9606"
/clone="IMAGE:2177941"
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Pred. No.
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                                                                          AI765236 456 bp mRNA linear EST 21-DEC-1999 wi72h08.xl NCI_CGAP_Kidl2 Homo sapiens cDNA clone IMAGE:3398911 3's similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                             GPR39.;, mRNA sequence.
AI765236
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TGGGCAGACACGAGCAGGAGAGACAG
                                                                               CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            High quality sequence stop: 'Location/Qualifiers
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Tumor Gene Index
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Seq primer: -40UP from Gibco
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a 172 c 143 g 121 t
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/tissue_type="2 pooled Wilms' tumors, one primary and one metastatic to brain"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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TTTGCCACAAGCAAGAGAATTTAACACTGTTTCAAACCCGGGGGAGTTGGCTGTTTA
                                                                                                                                                                        GAAAAACACAGCATGAGAAACACAGTAAAATGAATAAAAACCATAAAATATTTTAGCCCCTCTG 456
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                                         TTTGCCACAAGCAAGAGAATTTAACACTGTTTCAAACCCGGGGGAGTTGGCTGTTA 576
                                                                                                             TTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTTGGACTTGACAGCTTCT 516
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cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-blo.linl.gov/bbrp/image/image.html
Insert Length: 961 Std Error: 0.00
Seq Primer: -40UP from Gibco
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 456)
1 (bases 1 to 456)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
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Fatima Bonaldo. F
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/clone_lib="NCI_CGAP_Kid12"
/tissue_type="2 pooled tumors (clear cell type)"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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ACAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGG 476
                                   CAGCCATGATCCTCCGAATCTGGTTTGGGCATCCAGCATACGGCCAATGTCACAACAATCA 120
                                                                                                                                                          CAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCA 356
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NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological
Disorders and Stroke, Brain Tumor Genome Anatomy Project
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381 bp mRNA linear EST 16-OCT-2000 7171h01.yl NCI_CGAP_Brn20 Homo sapiens cDNA clone IMAGE:3340177 5' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CDNA Library Preparation: David B. Krizman, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL, send email to: info@image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1998)
Other_ESTs: 7171h01.x1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                              /note="Organ: brain; Vector: pAMP1; mRNA made from oligodendroglioma tissue, cDNA made by oligo-dT pr Directionally cloned. Size-selected on agarcse geaverage insert size 500 bp. Primary library, non-amplified."

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/clone_lib="NCI_CGAP_Brn20"
/tissue_type="oligodendroglioma"
/dev_stage="adult"
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/db_xref="taxon:9606"
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NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/nciagap.
NCI/NINDS-CGAP inttp://www.ncbi.nlm.nih.gov/nciagap.
National Cancer Institute / National Institute of Neurological
Disorders and Stroke, Brain Tumor Genome Anatomy Project
CCGAP/BTGAP), Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: Robert Jenkins, M.D., Mark Israel, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1998)
Other_ESTs: 7169f08.x1
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
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                                                                                                                                                                                               average insert size 500 bp.
non-amplified. "
81 c 93 g 83 t
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                                                                                                                                                                                                                                                                                                                             /dev_stage="adult
/lab_host="DH10B"
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/tissue_type="oligodendroglioma"
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/clone="IMAGE:3339975"
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ∕organism="Homo sapiens"
                                                                                                                       58.9%;
                                                                                                    score 368; DB; Pred. No. 0; 0; Mismatches
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                                                                                                                                            DB 12;
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578 bp mRNA linear EST 19-DEC-wg38h03.xl Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone IMAGE:2367413 3' similar to SW:GP39 HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR GPR39. ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Insert Length: 806
Seq primer: -40UP f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IMAGE Consortium (info@image.llnl.gov)
Insert Length: 806 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL ; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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            /note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from five normalized libraries were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 5 libraries. The pools consisted of the following libraries and cloneIDs: Soares NbHSF pool 1: 309384-310919, 333208-325895 Soares NbHSP pool 1: 145032-147335, 147720-148103, 148872-149255, 15002 - 150407, 151176-155327 Soares NbHSF-9w pool 1: 758280-760583, 772104-774407 Soares NbHRP pool 1: 733720-726407, 73980-740999 Subtraction by Bento pool 1: 733720-726407, 73980-740999 Subtraction by Bento
                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2367413"
/clone_lib="Soares_NSF_F8_9W_OT_PA_P_S1"
/lab_host="DH10B"
Soares and M. Fatima
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  Bonaldo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGGGCAGACACGAGCAGGAGGGAGAGAGAAAAGAAAAACACAGGCATGAGAAC 416
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                                                                                                                                                       Seq
                                                                                                                                                                                                                                                                                                               Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life
Technologies, Inc. cDNA Library Arrayed by: Christa Prange,
I.M.A.G.E. Consortium DNA Sequencing by: Washington Universi
                                                                                                                                                                                                                                                Genome Sequencing Center Clone distribution Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GPR39.;, mRNA sequence.
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                                                                                                                                                                              Possible reversed clone: similarity
                                                                                                                                                                                                                              www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tumor Gene Index
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National Cancer Institute, Cancer Genome Anat
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                                                                                                       quality sequence stop: 306 Location/Qualifiers
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                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
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TITLE
JOURNAL
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347; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mRNA sequence.
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/tissue_type="melanotic melanoma"
/lab_host="DH10B (phage-resistant)"
                                                /clone="IMAGE:4026092"
/clone_lib="NIH_MGC_20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               55.5%;
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560 GGGAGTTGGCTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG 606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           353 AAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        260 AAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            380 GGGAGAGACAGAAAAAGAAAAACACAGCATGAGAACACAGTAAATGAATAAAACCATAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGGACTTGACAGCTTCTTTTGCCACAAGCAAGAGAGAATTTAACACTGTTTCAAACCCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGT 499
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                                                                                                      cDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Colone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov Plate: LLCM855 row: o column: 21 High quality sequence stop: 723.
                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                  NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BF126050 759 bp mkNA linear L31 24 V 601762888F1 NIH_MGC_20 Homo sapiens CDNA clone IMAGE:4026092
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Site_2: Notl; Cloned unidirectionally. Primer: Oligo dT
Library constructéd by Life Technologies."
1 16 c 89 g 134 t 1 others
/organism="Homo sapiens"
/db_xref="taxon:9606"
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/tissue_type="2 pooled Wilms' tumors, one primary
metastatic to brain"
/lab_host="DH10B"
                                                                            Location/Qualifiers
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Pred. No. 0;
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RESULT 31
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mRNA sequence.
BF125134
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                                                                                                                             High quality sequence stop: 711.
Location/Qualifiers
                                                                                                                                                                            Plate: LLCM853 row: p column: 13
                                                                                                                                                                                            cDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.
                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1999)
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="InhAGE:4025340"
/clone_lib="NIH_MGC_20"
/tissue_type="melanotic melanoma"
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Pred. No.
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                This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Possible reversed clone: similarity on wrong strand Possible reversed clone: polyI not found Seg primer: -400P from Gibco
                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 387)

Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Krizman, D., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AI272281 387 bp mRNA linear EST 18-NOV-1998 ap23c03:x1 Schiller oligodendroglioma Homo saplens cDNA clone IMAGE:1956196 3' similar to SW:NTRL_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1 contains TAR1:t3 TAR1 repetitive element; mRNA sequence.
                                                                                                                                                    Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                Washington University Scho
4444 Forest Park Parkway,
                                                                                                                                                                                                                                           Unpublished (1997)
Contact: Wilson RK
                                                                                                                                  Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                       White,Y., Wylie,T., Waterston,R. WashU-NCI human EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EST
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High quality sequence stop: 386
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/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
ECORI; CDNA made by oligo-off priming. Directionally
cloned into EcoRI/XhoI sites using the following 5'
adaptor: GGCACGAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
a 192 c 255 g 171 t
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Pred. No.
                                                                                                                                                                                                  School of Medicine way, Box 8501, St. 1
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                                                                                                                                                 AW191974
x178a02.x1 NCI_CGAP_Pan1
similar to SW:GP39_HUMAN
Unpublished (1997)
Contact: Robert Strausberg,
                                        Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 454)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                                                                                    AW191974
AW191974.1 GI:6470673
                        Tumor Gene Index
                                                                                   Homo sapiens
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                                 National Cancer Institute,
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                                                                                                                                      ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /tissue_type="oligodendroglioma"
/dev_stage="44 years"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="Schiller oligodendroglioma"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
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Pred. No. 0;
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Homo sapiens cDNA clone IMAGE:2680778 3'
043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
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173 GAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAG 232
                                                                                                                                                                                                                AAGGGAAGAGATCATGTACGCCCGGAAGTAGGACCTCCTCCAGTCGTGCTTGGGTTTG
                                                                                                                                                                                                                                                                                             GAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGC 472
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    A1423162 480 bp
tf32c12.x1 NCI_CGAP_Brn23 Homo s
similar to SW:NTR1_RAT P20789 NE
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    www-bio.llnl.gov/bbrp/image/image.html
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/db_xref="taxon:9606"
/clone="IMAGE:2680778"
/clone_lib="NCI_CGAP_Pan1"
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/lab_host="DH10B"
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REFERENCE

ORGANISM

Homo sapiens

AUTHORS TITLE

JOURNAL

Contact: Robert Strausberg, Ph.D.

cgapbs-remail.nih.gov

Bonaldo, Ph.D.

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133 GCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAG 192
                        TCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAGAGAGAATTTAACACTGTTTCA
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                                                                                                                                                                                                                                                                                       AATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACG 372
                                                                                                                                                                                                                                                                                                                                                         CGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCG
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                                                                    ACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTT
                                                                                                                                                                 AGCAGGAGGGAGAGACAGAAAAAAAAACACAGCATGAGAACACAGTAAATGAATAAA 432
1 (bases 1 to 480)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NATIONAL Cancer Institute / National Institute of Neurola Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
Unpublished (1998)
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/lab_host="DH10B"
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                                                                                             ATCAGCCCTGGGCAGACACGAGCAGGAGGGGAGAGAGAAAAGAAAAACACAGCATGA 195
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R. and Wilson,R.
Washi,NCI human EST Project
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 314 286 1800
Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AA665640.1 GI:2620253
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1997)
                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                       78
                                                                                                                                                                                                                                                                                                                                                                                              /note="vector: pSPORT1; Site_1: Sall; Site_2: Not1; RNA was prepared from a pool of 6 anonymous Wilms' tumor RNAs. RNA was prepared by acid-phenol, followed by one round of oligo dT selection. cDNA library preparation was with the BRL/Life Tech. Superscript Plasmid System. An oligo-dT NotI primer for first strand synthesis generated geggegecc(t)n at the 3' end of the clones. A 5' Sall adaptor was used with sequence 5'-gtcgaccacgcgtccg-3'. Resulting cDNAs were size selected (average size 2 kb), NotI digested, and ligated into NotI/SalI-cut pSPORT1. Library was constructed by Dr. Manfred Gessler."
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/sex="pooled (6)"
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/db_xref="taxon:9606"
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quality sequence stop: 372.

Location/Qualifiers

/clone="IMAGE: 2097910" /clone\_lib="NCI\_CGAP\_Brn23"

/db\_xref="taxon:9606" /organism≃"Homo sapiens"

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                  GAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGC 472
                                                                                                                         GCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACA 352
                                                                                GAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGC 136
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tumor Gene Index Unpublished (1997)
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1 (bases 1 to 354)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                          /note-"Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687239, 726408-728711, and 729096-731399. Subtraction by Bento Soares and M. Fatima Bonaldo."
                                                                                                                                                                                                                                                                         Soares and M. Fatima Bonaldo.
88 c 78 g 108 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="Soares_NFL_T_GBC_S1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="IMAGE:1580506"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ocation/Qualifiers
                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                              0;
                                                                                                                                                                                                          Score 314; DB 9; Length 354; Pred. No. 0;
                                                                                                                                                                                          Mismatches
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AA075710/c
                                              BASE COUNT
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ORGANISM
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Query Match
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                                                                                                                                                                                                                                                                                     source
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593 TGCTTTAGACAGTG 606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: estewatson.wustl.edu WARNING: There is evidence that suggests that the 384-well parent plate of this clone contains both human and mouse derived clones. Thus, the origin of this clone is uncertain. This caution should kept in mind should you use this clone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 373)
Hillier,L., Lennon,G., Becker,M., Bonaldo,M.F., Chiapelli,B.,
Chissoe,S., Dietrich,N., DubBuque,T., Favello,A., Gish,W., Hawkins
,M., Hultman,M., Kucaba,T., Lacy,M., Le,M., Le,N., Mardis,E., Moore
,B., Morris,M., Parsons,J., Prange,C., Rifkin,L., Rohlfing,T.,
Schellenberg,K., Soares,M.B., Tan,F., Thierry,Meg,J., Trevaskis,E.,
Underwood,K., Wohldmann,P., Waterston,R., Wilson,R. and Marra,M.
Generation and analysis of 280,000 human expressed sequence tags
Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 427 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA075710
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97044478
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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h quality sequence stop: 240.
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314 286 1810
                                                                       83
                                                               /organism="Homo sapiens"
/db_xref="GDB:3923283"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                          /clone_lib="Stratagene ovarian cancer (#937219)"
                                                                                                                                                                                                                                                                                                                                         /clone="IMAGE:545095"
                                                                                                                                                                                                                                                                                           /sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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Score 314;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             373 bp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          412 bp mRNA linear EST 31-JUL-19. zn17f09.s1 Stratagene neuroepithelium NT2RAMI 937234 Homo sapiens cDNA clone IMAGE:547721 3', mRNA sequence.
                                                                                                                                                                                                                                                                                         This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 980 Std Error: 0.00 Seq primer: -40M13 fwd. from Amersham.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
/clone_lib="Stratagene neuroepithelium NT2RAMI 937234"
/dev_stage="Ntera-2/RA+MI neuroepithelial cells"
/lab_host="SOLR (kanamycin resistant)"
/note="Vector: pBluescript SK-; Site_1: EcoRI; Site_2: XhoI; Cloned unidirectionally. Primer: Oligo dT. NT2
(Ntera-2/c1.D1) precursor cells induced with Retinoic Acid for 1 week, followed by 3 weeks in mitotic inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hominidae;
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Tel: 301 402 3452
Fax: 301 496 0078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         by09g04.yl Human Lens cDNA (Un-normalized, unamplified): sapiens cDNA clone by09g04 5', mRNA sequence.

BF726644
                                                                                                                                                                                                                                                                                       Email: graeme@helix.nih.gov
Plate: 09 row: g column: 04
Seg primer: M13RP1 reverse primer (ABI).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wistow,G.J., Bernstein,S., Behal,A. and Smith,D. NEIBANK: EST analysis and bioinformatics for ocular Invest. Ophthalmol. Vis. Sci. 41, (2000) In press
                                                                                                                                                                                                                                                                                                                                                                                                                      Section on Molecular Structure and Function National Eye Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wistow G
  /note="Organ: Eye; Vector: pCMVSPORT6; Two human lenses from different adults (both approximately 40 years old) together yielded 20ug of total RNA and 150ng mRNA for cl
                                                                    /dev_stage="Adult"
/lab_host="EMDH10B"
                                                                                                                                                                              /clone="by09g04"
                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                            /tissue_type="Lens"
                                                                                                                                                            /clone_lib="Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            606
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                                                                                                                                                            cDNA (Un-normalized, unamplified):
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FEATURES

Location/Qualifiers /organism="Homo sapiens"

source

/db\_xref="GDB:3926237" /db\_xref="taxon:9606" /clone="IMAGE:547721"

COMMENT

Contact: Wilson RK
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. I
Tel: 314 286 1800
Fax: 314 286 1810
Email: est@watson.wustl.edu

St. Louis,

TITLE JOURNAL MEDLINE

REFERENCE

(bases 1 to 412)

AUTHORS

SOURCE VERSION ACCESSION KEYWORDS

ORGANISM

Homo sapiens human AA084249.1

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(Replate #2). Average insert size: 1.1 kb; Uni-ZAP Vector; ~5' adaptor sequence: 5' GAATTCGGCACGAG 3' adaptor sequence: 5' CTCGAGTTTTTTTTTTTTTTTT 3'" 110 c 87 g 126 t
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REFERENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA173739
zp04h05.r1 Stratagene ovarian
clone IMAGE:595449 5', mRNA se
                                             Email: est@watson wustl.edu
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Insert Length: 792 Std Error: 0.00
                                                                                                                                                                                            Unpublished (1997)
Contact: Wilson RK
Washington University School of Medicine
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis,
                                                                                                                                                                                                                                                                                          Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Wartin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylle,T., Waterston,R. and Wilson,R. WashU-NCI human EST Project
                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 349)
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AA173739.1 GI:1753890
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quality sequence stop: 236
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        library synthesis. A directionally cloned cDNA library in the pcMVSPORT6 vector was constructed at Life Technologies , essentially following the protocols of the SuperScript Plasmid System full details of which are contained in the
                         -28M13 rev2 from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              49.9%; Score 312; 100.0%; Pred. No.
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                           Std Error: 0.00 for ev2 from ...
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cancer (#937219) Homo
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                                                                                                                                                                                                                                                                                                            408 bp mRNA linear EST 25-J
BG057775 Lupski_dorsal_root.ganglion Homo sapiens cDNA clc
7f75g10.x1 Lupski_dorsal_root.ganglion Homo sapiens cDNA clc
IMAGE:3302971 3' similar to SW:GP39 HUMAN 043194 PUTATIVE G
PROTEIN-COUPLED RECEPTOR GPR39. ; mRNA sequence.
                                                                       Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. James Lupski
                                                                                                                                                                                                                                                                   EST
                                                                                                                         Unpublished (1997)
                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                                                                                                                                                                                                                                                BG057775.1 GI:12523767
found through the I.M.A.G.E. Consortium/LLNL
        cDNA Library Preparation: Lupski Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Washington University Genome Sequencin
Clone distribution: NCI-GAP clone distribution informati
                                                                                                                                      Tumor Gene Index
                                                                                                                                                      National Cancer Institute, Cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="Stratagene ovarian cancer (#937219)"
/sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /dev_stage="adult, 64 years"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="IMAGE:595449"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /db_xref="taxon:9606"
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100.08;
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Pred. No.
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                                                                                                                                                     Genome Anatomy Project (CGAP),
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                             Sequencing Center
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FEATURES

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primer: -40UP f

source

/sex="male"

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GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA 479
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             1 (bases 1 to 420)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of N
Disorders and Stroke, Brain Tumor Genome Anatomy Pr
                                                                                                                                                                                                                                                   AI566797
420 bp mRNA linear EST 12-MAY-1999 tn23g08.xl NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2168510 3' similar to SW:NTR1_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1 ;, mRNA
                                                                                               Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       quality sequence stop:
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5'-GACTAGTTCTAGATCGCGAGCGGCCCT(15)-3'. Size selected 1 kb for average insert length 1.7 kb. This is a primary library, non-amplified. Library constructed by Life Technologies and donated by J. Lupski, M.D./Ph.D. (Baylc College of Medicine) and is available through Life Technologies."

79 g 128 t 3 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note-"Vector: pCMV-SPORT6 (Life Technologies); Site_1: Not1; Site_1: Sal1; cDNA made by oligo-dT priming. Directionally cloned using the following adaptors: 5'-TCGACCCACGCGTCCG-3' and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="dorsal root ganglia"
/dev_stage="adult, 36 yr"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:3302971"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             348;
                                                       qy56f02.x1
similar to
sequence.
AI363261
AI363261.1
                                                                                             AI363261
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: POLYA=No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bonaldo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tissue Procurement: David N. Louis, M.D., Myrna R. \text{Ph.D.}
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note-*organ: brain; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Solte_2: Eco RI; 1st strand cDNA was primed with a Not I: Site_2: Eco RI; 1st strand cDNA was primed with a Not I oligo(dT) primer [5]; Gotthe-stranded cDNA was ligated to Eco RI 3']; double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized, and was constructed by Bento soares and M.Fatina Bonaldo." 1 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="anaplastic oligodendroglioma"
/lab_host="DH10B"
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/clone_lib="NCI_CGAP_Brn25"
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/db_xref="taxon:9606"
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SW:NTR1_RAT P20789 I
 GI:4114882
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99.78;
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Pred. No. 0;
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                                                     bp mRNA linear EST 16-FEB-1999
o sapiens cDNA clone IMAGE: 2016027 3'
NEUROTENSIN RECEPTOR TYPE 1 ;, mRNA
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Query Match Best Local

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SOURCE ORGANISM VERSION KEYWORDS

Homo sapiens

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AAACACAGGCATGAGAACACAGTAAATGAATAAAACCATAAAATATTTTAGGCCCCTGTGTTC 459
                                                                                                                                                                         AAACGTCTCCGAGAAGGGGGGGGGGGTCATGTACGCCCGGAAGTAGGACCTCGTCCAGTC
                                 TGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTGGACTTGACAGCTTCTTTT 519
                                                                    AAACCCAGCATGAGAACCCCAGTAAATAAATAAAACCATAAAATATTTAGCCCCCTCTGTTC
                                                                                                                                      CAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAGGAGAGACAGAAAAGAA
                                                                                                                                                                                                            GTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGC
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CDNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 814 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 475)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq primer: -40UP from Gibco
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/clone="IMAGE:2016027"
/clone_lib="NCI_CGAP_Brn23"
/tlssue_type="glioblastoma (pooled)"
/lab_host="DH10B"
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99.38;
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Pred. No. 0;
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                                                                                                                                                                                                                                                                 GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
                               CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAAACGTCTCCGAGAAGGGGGAG
                                                                                                                                                                                                                                 GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                 CAGCGACAGGCGCAGCACCAGCACCAGGAACACCCGGCGAAACTGCTGCGAGGACAC
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found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
Plate: L/CM2388 row: o column: 21
High quality sequence stop: 665.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1999)
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NIH-MGC http://mgc.nci.nih.gov/.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             вуову//1 888 bp mRNA linear ES AGENCOURT_8344092 NIH_MGC_110 Homo sapiens cDNA clone 5', mRNA sequence.
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia; Eutheria;
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/clone_lib="NIH_MGC_110"
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/tlsb_host="DH1DB (phage-resistant)"
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/db_xref="taxon:9606"
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100.0%; Pred. No. 0;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                            344 TTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAAT 285
                                                                                                                                                                                                                                                                                                                                   284 TTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCCAGCATACGGCCAAT 343
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                                      ACAGCATGAGAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTG 463
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CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: MCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 402) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Possible reversed clone: similarity on wrong strand Seq primer: -40UP from Gibco High quality sequence stop: 303.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        www-bio.llnl.gov/bbrp/image/image.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="organ: ovary; Vector: pcMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.35 kb. Tumor types include: mixed Mullerian tumor, papillary serous, clear cell, spindle cell. All are primary tumors, metastasis positive. Life Technologies catalog #: 11534-013"

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/clone_lib="NCI_GGAP_Ov23"
/tlssue_type="tunor, 5 pooled (see description)"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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99.7%;
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Maximum DB
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Match Length
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1: /cgn2_6/ptodata/2/pubpna.
2: /cgn2_6/ptodata/2/pubpna.
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 /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
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US-09-825-294-205
US-09-825-294-211
US-09-867-701-4240
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US-09-867-701-8394
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4240, Ap
208, App
1516, Ap
1516, Ap
2409, Ap
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US-10052-545-11	US-09-815-242-4132	US-09-960-352-12765	US-09-969-347-117	US-09-148-012-1	US-09-804-551B-41	US-09-880-107-2251	US-09-964-824A-289	US-09-764-877-3803	US-09-764-869-2001	969	US-09-925-302-122	US-09-888-615-26	US-09-885-441-1	US-09-962-832-105	US-09-753-831-3	US-09-925-301-541	US-09-867-701-9973	US-09-993-844-14	-09-7		US-09-764-847-1091	US-09-764-847-1092	US-10-037-616-1	US-09-971-228-4	US-09-971-228-1
	Sequence 4132, Ap	Sequence 12765, A	Sequence 117, App	Sequence 1, Appli	Sequence 41, Appl	Sequence 2251, Ap	Sequence 289, App	Sequence 3803, Ap	Sequence 2001, Ap	Sequence 1999, Ap	Sequence 122, App	Sequence 26, Appl	Sequence 1, Appli	Sequence 105, App	Sequence 3, Appli	Sequence 541, App	Sequence 9973, Ap	Sequence 14, Appl	325	3251,	Sequence 1091, Ap	Sequence 1092, Ap	Sequence 1, Appli	Sequence 4, Appli	Sequence 1, Appli

## ALIGNMENTS

```
APPLICANT: M. Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Hing, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN C.
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 210
LENGTH: 625
TYPE: DNA
TYPE: DNA
TYPE: DNA
                                                                                                                                                                                                                                                                  FEATURE:

NAME/KEY: misc_feature
LOCATION: (1)...(625)
OTHER INFORMATION: n = A,T,C
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                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 625; Conserv
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121
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              CAGCGACAGGCGGCAGCACACCTGCACGAACACCCGCCGAAACTGCTGCGGAGGACAC 180
                                                              CAGCGACAGGCGGCAGCACCAGCACCGAACACCCGGCGAAACTGCTGCGAGGACAC
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Sequence

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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Jones, Robert
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 10876
LENGTH: 625
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                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens FEATURE:
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                                                              GCCCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                 AGTTCTCCTTGCAGAGGACTGGCGCGGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
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100.0%; Pr
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                                                                                                                                                    Score 624; DB 10;
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Sequence 214, Application US/09825294
Patent NO. US20020004491A1
GENERAL INFORMATION:
APPLICANT: XL, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
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                                                                                                                                                                                  ; NAME/KEY: misc_feature
; LOCATION: (1)...(1897)
; OTHER INFORMATION: n =
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 214
LENGTH: 1897
TYPE: FOR
                                                                                                                                Matches 624;
                                                                                                                                                          Query Match
                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: HOMO :
FEATURE:
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                                       GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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                         GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                             AGTTCTCCTTGCAGAGGACTGGCGCCGGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
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APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Filing, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 205
LENGTH: 1619
TYPE: DNA
ORGANISM: Homo sapiens
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CAGCGACAGGCGCAGCACAGCACC-TGCACGAACACCCGGCGAAACTGCTGCGAGGACA 179
                                                                     GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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                                                     GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 1111
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99.7%;
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SOFTWARE: FASTSEQ FOR WINDOWS VERSION SEQ ID NO 211
LENGTH: 1619
TYPE: DNA
ORGANICO:
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                                                                                                                                                                                                                        Matches 624;
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Best Local (
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APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHOD
TITLE OF INVENTION: THERAPY AND DIAGNOSIS
FILE REFERENCE: 210121.484C5
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                          CAGCGACAGGCGGCAGCACAGCACC-TGCACGAACACCCGCCGAAACTGCTGCGAGGACA 179
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                                                                                                                        GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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Pred. No. 5e-156;
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US-09-867-701-2375

Sequence 2375, Application Patent No. US20020132237A1 GENERAL INFORMATION:

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APPLICANT: Jones, Robert

APPLICANT: Jones, Robert

APPLICANT: Jones, Robert

APPLICANT: Jones, Robert

APPLICANT: Herlocker Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF ESO ID NOS: 10912

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 4240

LENGTH: 409

TYPE: DNA

ORGANISM: Homo sapien
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                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 407; Conserv
  170
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                                                                                                                                                                                                           198 TGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCC
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AAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGT
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                                                              GAGGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAGTAAATGAATAAAACCAT
                                                                                                          GGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAG
                                                                                                                                                                                               TGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCC
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                                                 GGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAG
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99.5%;
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                                                                                                                                                                                                                                                        Score 395.4; DB 1
Pred. No. 1.4e-97;
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APPLICANT: AGIACE, Paul A.

APPLICANT: Jones, Robert
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2375
LENGTH: 558
TYPEN: DNA
CAGCGACAGGCGGCAGCACCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCCCTCTCGCTGGAGTGCGCATGTACGCG--AGGCCTTCTCGTGGTTGGCGTGCTG
                                                                                                                           GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                     AGTTCTCCTTGCAGAGGACTGGCGCGGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
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                                                                                                                                                                                                                                                     Score 433.2; DB 10;
Pred. No. 1e-107;
0; Mismatches 14;
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\_  $\vdash$  Query Match Best Local S Matches 539

Similarity

69.3%;

Conservative

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; OTHER INFORMATION: n - A,T,C US-09-867-701-2375

or,

NAME/KEY: misc\_feature LOCATION: (1)...(558)

ORGANISM: Homo sapien

FEATURE:

TYPE: DNA

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119

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1:

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NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 208
LENGTH: 1362
TYPE: DNA
ORGANISM: Homo sapiens
                                                      RESULT 9
US-09-867-701-1516/c
Sequence 1516, Application US/09867701
Patent No. US20020132237A1
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                                              GENERAL INFORMATION:
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Best Local Similarity 98.9%;
Matches 364; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
           APPLICANT: Aglate, Paul A. APPLICANT: Jones, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Xu, Jiangchun APPLICANT: Stolk, John 1
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 Harlocker,
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Pred. No. 3.5e-88;
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APPLICANT: Jones, Robert
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1532
LENGTH: 390
TYPE: DNA
ORGANISM: Homo sapien
                                         ; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(390)
; OTHER INFORMATION: n =
US-09-867-701-1532
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CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1516
LENGTH: 373
                                                                                                                                                                                                                                                                                                                              Sequence 1532, Application US/09867701 Patent No. US20020132237A1
                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             594 GCTTTAGACAGTG 606
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99.7%;
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Score
Pred.
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Pred. No. 2.6e-88;
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353; DB 10;
No. 3.8e-86;
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             Length 390;
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Matches

278 61

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Sequence 2409, Application US/09867701

Patent No. US20020132237A1

GENERAL INFORMATION:
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT EILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FRASTSEQ for Windows Version 4.0
SEQ ID NO 2409
TYDE: DAM
                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-867-701-2409
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US-09-867-701-2409/c
                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Sim
Matches 328;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapien
                                                                                                                                                                                                                   274 CCAGTCGTGCTTGGGTTTGGCCGC---AGCCATGATCCTCCGAATCTGGTTGGGCATCCA 330
                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
GCTTCTTTTGCCAAGCAAGAGAGAGATTTAACACTGTTTCAAACCCGGGGGAGTTGGCT 570
                                    CCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTGGACTTGACA 510
                                                                                                                                     AGAAAAGAAAAACACAGGATGAGAACACAGTAAATGAATAAAACCATAAAATATTTAGCC
                                                                                                                                                                                                                                                                            CCAGTCGTGCTTGGGGTTTTGCCCGCAGCCCATGATCCTCCGAATCTGGTTGGGCATCCA 280
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                                                                                                                    AGAAAAGAAAACACAGCATGAGAACACAGTAAATGAATAAAACCATAAAATATTTAGCC
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96.8%;
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Pred. No. 8.9e-75;
"" matches 8;
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CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7826
LENGTH: 381
TYPE: DNA
ORGANISM: Homo sapien
US-09-867-701-7826
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                                                           US-09-867-701-8894
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                                                                     CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FASESEQ FOR WINDOWS Version 4.0
SEQ ID NO 8894
LENGTH: 201
TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                    Sequence 8894, Application US/09867701 Patent No. US20020132237A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 7826, Application US/09867701 Patent No. US20020132237A1
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                Query Match
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Best Local
Best Local Similarity
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APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
                                                                                                                                                                                                                             APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Aglate, Paul A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               328 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAG-AAAACGTCTCCGAGAAG 381
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Score
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171.2; DB 1
No. 3.3e-37;
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TYPE: DNA

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Sequence 17, Application US/09866582
Patent NO. US20020127620A1
GENERAL INFORMATION:
APPLICANT: Witman, George B.
APPLICANT: Pazour, Gregory J.
APPLICANT: Rosenbaum, Joel L.
APPLICANT: Cole, Douglas G.
TITLE OF INVENTION: INTRAFLAGELLAR TRANSPORT
FILE REFERENCE: 07917-145001
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US-09-866-582-17/c
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US-09-149-045-1
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CURRENT APPLICATION NUMBER: US/09/149,045

CURRENT FILING DATE: 1998-09-08

NUMBER OF SEQ ID NOS: 8

SOFTWARE: PATENTIN VET. 2.0 - beta

SEQ ID NO 1

LENGTH: 1464

TYPE: NAME
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Best Local Similarity 52.6%;
Matches 123; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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APPLICANT: HORUK, RICHARD
APPLICANT: ZHOU, YIQING
APPLICANT: HESSELGESSER, JOE
APPLICANT: DOMS, ROBERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1182 CTGGCCACAGCAGCATGGAGGTGCAGGCCTGGCGGAAGCGGGGGTCGAAAAAGGCCATA 1123
                                                                                                                                                                                                                                                                                                                                               1062 CATGAGGAAGAGGTCAAAGTCACAGGGCCAGTGCAGCAGGCTGCCCAGCATGTACAGCGT
                                                                                                                                                                                                                                                                                                                                                                                                                                 1112 GAGGAAGGGGTTGAGCAGCTGTTGACGTAGCTGATGCAGGTGCAGTAGGGGAAGATGTT
                                                                                                                                                                                                                                                                    1002 CTTCACCAGGTGGTAGGGCATCCAGCACAGGGCAAAGGTCACCACCAGCACCAC 949
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       127 CAGGCGGCAGCACCAGCACCAGCACGCCGCCGAAACTGCTGCGAGGACACCGTGTA 186
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   231 AGAAGGGAGGAGCATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTCCTTGGGTT 290
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; LOCATION: (3)...(1517)
US-09-866-582-17
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17
LENGTH: 2036
                                                                                                                                                                                                                                                                                                                                   Best Local Similarity Matches 103; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/866,582
CURRENT FILING DATE: 2001-05-24
PRIOR APPLICATION NUMBER: US 60/206,923
PRIOR FILING DATE: 2000-05-24
NUMBER OF SEQ ID NOS: 50
time : 34.1619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Chlamydomonas reinhardtii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                362 CTTGATGCTCTTGAG 348
                                                                                                                               422 CAGCTCCGTGAAGCGCTCCAGCGCGGAGCGGTCCGAGGGCGCCATGTTGGAGGGCGGGTT
                                                                                                                                                   139 CAGCACCTGCACGAACACCCGCCGAAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTT 198
                                                                                                                                                                                                482 GGGCGAGTGCACGGACTTGTGCACCACTTCGTAGGCGTAGTACACCTCGGCGCGGTCGTA
                                                                                                                                                                                                                                                               199 GATGACCGAGCTGAG 213
               completed: November
                                                                                                                                                                                                                              79 GTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCA 138
                                                                                                                                                                                                                                                                                               19 CTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCGGGGGGCGCTGTCGGTGGTA 78
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secs
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              2002, 18:00:05

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Pred. No. 0.0023;
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and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1.
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
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/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
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US-09-299-843A-39
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US-09-088-337B-39
PCT-US93-11153-39
US-08-232-463-14
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## ALIGNMENTS

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RESULT 1
US-08-153-848-39/c
^-~~once 39, Application US/08153848
US-08-153-848-39
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                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/153,848
FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/977,452
FILING DATE: 17-NOV-1992
ATTORNEY/AGENT INFORMATION:
NAME: NO. 5759804and, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 31794
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                             TELEPHONE: (312) 474-630
TELEFAX: (312) 474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: :
                FEATURE:
NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Godiska, Ronald
APPLICANT: Gray, Patrick W.
APPLICANT: Schweikart, Vicki L.
TITLE OF INVENTION: NO. 5759804el Seven Transmembrane Receptors
NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 1872 base pairs
                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
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CITY: Chicago
STATE: Illinois
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                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                              FILING DATE: 17-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/977,452
FILING DATE: 17-NOV-1992
ATTORNEY/AGENT INFORMATION:
TILL F. Uhl
MOLECULE TYPE: FEATURE:
                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                           TELEPHONE: (312) 474-6300
                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 09/088,337 FILING DATE: 01-JUN-1998 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Godiska, Ronald
APPLICANT: Gray, Patrick W.
APPLICANT: Schweikart, Vicki L.
TITLE OF INVENTION: No. 6107475el Seven Transmembrane Receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
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                                    TOPOLOGY:
                                                                       TYPE:
                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 27
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STATE: Illino
                                                   STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 0
FILING DATE: 17-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 39, Application US/09088337B Patent No. 6348574
GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches 123;
                                                                                                                                      TELEFAX: (312) 474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 39:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1005
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                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/153,848
FILING DATE: 17-NOV-1993
APPLICATION NUMBER: US 07/977,452
FILING DATE: 17-NOV-1992
ATTORNEY/AGENT INFORMATION:
NAME: NO. 6348574and, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 31794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   247 CATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGAT 306
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/088,337B
FILING DATE: 01-Jun-1998
                MOLECULE
                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: NO. NUMBER OF SEQUENCES: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Godiska, Ronald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAAATCAACCAATCAGCCC 360
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                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                LENGTH: 1872 base pairs
                                                                                                                                                                                                TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 6300 Sears Tower, 233 South Wacker Drive CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Marshall, O'Toole, Gerstein, Murray &
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                TYPE: DNA (genomic)
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Schweikart, Vicki L.
                                   linear
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                                                                                                                                                                                                    (312) 474-6300
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                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/977,452
FILING DATE: 17-NOV-1992
ATTORNEY/AGENT INFORMATION:
NAME: NOLand, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 31794
                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1872 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1005 CTTCACCAGGTAGGGCATCCAGCACAGGGCAAAGGTCACCACCAGCACCAC 952
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                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole, Gerstein,
ADDRESSEE: Bicknell
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                                                                                                                                                                                               TELEPHONE: (312) 474-6300
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MEDIUM TYPE: Floppy disk
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                                  MOLECULE TYPE: DNA (genomic)
                                                                                                                                             TELEFAX: (January 1975)
TELEFAX: 25-3856
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Schweikart, Vicki L.
TITLE OF INVENTION: Novel Seven Transmembrane Receptors
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 6300 a
                                                                LENGTH: 1872 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE:
NAME/KEY:
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                                                     TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                               (312) 474-0448
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Version #1.25
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US-08-499-215-1/c
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                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: JP H
FILING DATE: 08-JUL-1994
INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, V SOFTWARE: & WordPerfect version 5. CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: DC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                           HYPOTHETICAL: NO ORIGINAL SOURCE:
                                                                                                                                                                                                            TOPOLOGIE TYPE: DN
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                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 6379 base pair
                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1005 CTTCACCAGGTGGTAGGGCATCCAGCACAGGGCAAAGGTCACCACCAGCACCAC 952
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FEATURE:
                                                                     FEATURE:
                                                                                                                                          FEATURE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: BIOLOGICAL DEGRADATIVE
TITLE OF INVENTION: TREATMENT OF CHLORINE-SUBSTITUTED ETHYLENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Saeki, Hisas
APPLICANT: Miura, Akira
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                               NAME/KEY:
LOCATION:
                                                                                  NAME/KEY: CDS
LOCATION: 910..1935
OTHER INFORMATION: ,
                                                                                                                                                           STRAIN:
                                                                                                                                                                                                                                                                  STRANDEDNESS:
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CITY: N.W.
                OTHER INFORMATION:
                                                                                                                                                                              STRAIN:
                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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                                                                                                                                                                                                                                                            nucleic acid
pedness: single
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                                                                                                                                                         No. 5612204ardia corallina B-276 (FERM P-4094; FERM BP-5124; ATCC 31338)
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2100 Pennsylvania
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Avenue
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                                                                                      "amoA'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Version #1.30
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AGCCATGATC 307

1293

238

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1233 CCCCATGGTC 1224

US-08-232-463-14/c

Sequence 14, Application US/08232463 Patent No. 5670367

APPLICANT: DORNER, F.
APPLICANT: SCHELFLINGER, F.
APPLICANT: FALKMER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLFOX VIRUS
NUMBER OF SEQUENCES: 52

CORRESPONDENCE ADDRESS:

STREET:

1800 Diagonal Road,

Suite 500

Foley & Lardner

Alexandria ٧A

ADDRESSEE:

GENERAL

INFORMATION:

APPLICATION NUMBER: EP 91 114
FILING DATE: 26-AUG-1991
ATTORNEY\_AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 3047
TELECOMMUNICATION INFORMATION:

EP 91 114 300.6 US/07/935,313

TELEFAX: TELEPHONE:

(703)683-4109

(703)836-9300

30472/114 IMMU

APPLICATION NUMBER: US FILING DATE: CLASSIFICATION: 435 PRIOR APPLICATION DATA: APPLICATION NUMBER: US

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:

Version

US/08/232,463

COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk

22313-0299

COUNTRY:

USA

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118

밁 δÃ ; NAME/KEY: CDS ; LOCATION: 3805..4830 ; OTHER INFORMATION: /I US-08-499-215-1

/product= "amoD"

FEATURE:

LOCATION: 2300..38 OTHER INFORMATION:

2300..3802

/product= "amoC"

NAME/KEY:

Query Match Best Local Similarity

8.0%;

Score 50; DB 1; Pred. No. 0.00054;

Matches

125;

Conservative

0;

Mismatches

125;

Вр

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1413 GTCCTGGAGCAGCCGCATGCGGTCCACGGCCTGGAACACCACGCTGAACTGGACCGTGTC 1354
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                                                                                                                                                                                                                                                           APPLICANT: Strand, Joann
APPLICANT: Hadcock, John R.
APPLICANN: Hadcock, John R.
TITLE OF INVENTION: No. 6420137el cDNA Encoding The Long and
TITLE OF INVENTION: Forms of The Human Neurotensin Subtype 2
FILE REFERENCE: 0630/0E792
CURRENT APPLICATION NUMBER: US/09/200,090
CURRENT FILING DATE: 1998-11-25
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 3
LENGTH: 1176
TYPEE. NA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence 3, Application US/09200090
Patent No. 6420137
                                                                                                                                   Matches 124;
                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                               TYPE: DNA ORGANISM: human
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IMMEDIATE SOURCE:
CLONE: pTZgpt-
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                                                                                            111 TGGCGTGCTGCAGCGACAGGCGGCAGCACCAGCACCTGCACGAACCCCGCCGAAACTGCT 170
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171 GCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCG
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                                                        TGGGGTGGTGCTCTCCACACAGGGAGCTGACGGCTTCCAGGAAGAGTTTTCTGAAGGAGG 1031
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                                                                                                                                   Conservative
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Pred. No. 0.00083;
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                                                                                                                                                     Score 46.6; DB Pred. No. 0.002;
                                                                                                                                   Mismatches 129;
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                                                                                                                                                                    DB 4;
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CURRENT APPLICATION NUMBER: US/09/007,005B
CURRENT FILING DATE: 1998-01-14
EARLIER APPLICATION NUMBER: 60/035,963
EARLIER APPLICATION NUMBER: 60/064,491
EARLIER APPLICATION NUMBER: 60/064,491
EARLIER FILING DATE: 1997-11-06
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17
EARLIER FILING DATE: 1997-11-06
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RESULT 9
US-09-244-796-17
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APPLICANT: SZOSTAK, Jack W.
APPLICANT: ROBERTS, RICHARD W.
APPLICANT: Liu, Rihe
TITLE OF INVENTION: SELECTION OF PROTEINS USING RNA-PROTEIN
TITLE OF INVENTION: FUSIONS
FILE REFERENCE: 00786/350003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17, Appl: Patent No. 625858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: (1)...(289)
OTHER INFORMATION: n = A,T,C or G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA
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                                                                                                      AACACA 406
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                                                                                                                                                              TGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCC
                                                                                                                                                                                                                                                                                                                              NRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          th 7.4%; Score 46; DB 4; Length 289, Similarity 9.3%; Pred. No. 0.0015; Similarity 103; Mismatches 120; Indels
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. LUCATION: (1)...(289); OTHER INFORMATION: n = A,T,C
US-09-244-796-17
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                                                                                                                                                                                                                                                           Sequence 1, Application Patent No. 6008050 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 23; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: SELECTION OF PROTEINS USING RNA-PROTEIN TITLE OF INVENTION: FUSIONS FILE REFERENCE: 00786/350007

CURRENT APPLICATION NUMBER: US/09/244,796

CURRENT FILING DATE: 1999-02-05

EARLIER APPLICATION NUMBER: 60/35,963

EARLIER FILING DATE: 1997-01-27

EARLIER APPLICATION NUMBER: 60/064,491

EARLIER APPLICATION NUMBER: 60/064,491

EARLIER FILING DATE: 1997-11-06

EARLIER FILING DATE: 1997-11-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17, Application US/09244796 Patent No. 6281344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Szostak, Jack W. APPLICANT: Roberts, Richard W. APPLICANT: Liu, Rihe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EARLIER APPLICATION NUMBER: 09/007,005
EARLIER FILING DATE: 1998-01-14
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 289
TYPE: RNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Translation template
                                                                                                                      NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Smithkline Beecham Corporation
COMPUTER READABLE FORM
                                                                                                                                                                                   APPLICANT: Shabon, Usman TITLE OF INVENTION: NOVET
                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            156
                                                                                                                                                                                                                                                                                                                                                                                                                                                            401 AACACA 406
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                                                                                                                                                                                                                                                                                                                                                                                                                      276 AAAAAA 281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        221 AACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCG 280
                                                               STREET: 709 Swedeland CITY: King of Prussia STATE: PA
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                        ZIP:
                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 36 CRARARURURARCRARARURGRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNR 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNR 155
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCC 340
                      19406
                                                                                                                                                                                                                                       Bergsma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7.4%; Score 46; DB 4; Length 289 ilarity 9.3%; Pred. No. 0.0015; Conservative 103; Mismatches 120; Indels
                                                                                                                                                                                              NOVEL HUMAN NEUROTENSIN RECEPTOR TYPE
                                                                                                                                                                                                                                                                                                         US/08832399
                                                                                                          Road
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TYPE:

US-09-372-498-1

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; MOLECULE TYPE: CDNA US-08-832-399-1
                                                                                               GENERAL INFORMATION:
APPLICANT: Derk J. Bergsma
APPLICANT: Usman Shabon
TITLE OF INVENTION: NOVEL HUMAN NEUROTENSIN RECEPTOR TYPE
TITLE OF INVENTION: AND SPLICE VARIANTS THEROF
FILE REFERENCE: GH-50020-1
CURRENT APPLICATION NUMBER: US/09/372,498
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: 08/832,399
PRIOR FILING DATE: 1997-04-02
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/09372498 Patent No. 6166182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1342 base pairs
  TYPE: DNA
ORGANISM: HOMO SAPIENS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: King, William T
REGISTRATION NUMBER: 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: GH
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      926 TGGGGTGGTGCTCTCCACACAGGGAGCTGACGGCTTCCAGGAAGAGTTTTCTGAAGGAGG 867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         111 TGGCGTGCTGCAGCGACAGGCGCAGCACCAGCACCTGCACGAACACCCGGCGAAACTGCT 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/832,399 FILING DATE: 02-APR-1997 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM:
SOFTWARE: FastSE(
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mes 120; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAA 350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TCACCATGTAGAAGTAGTAGAAATTGTACAGTGGGTCAGTCCACGCGTCATCAGGTA 747
                                                                               FastSEQ for Windows Version 3.0
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Best Local Similarity 49.2
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                                                                                                                                                                                                                    NAME: Player, William E.
REGISTRATION NUMBER: 31,049
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Danies .....APPLICANT: Pascale CHALON APPLICANT: Pascual FERRARA
                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 19-SEP-19:
                                       FEATURE:
                                                                               MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 1575 base pairs
                                                                                                                                                                                                                                                                            APPLICATION NUMBER: PCT/
FILING DATE: 17-MAR-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 536 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 20004
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Pascual FERR APPLICANT: Vita NATALIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    171 GCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCG
NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          686 CGAT 683
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  111 TGGCGTGCTGCAGCGACAGGCGGCAGCACCAGCACCAGCAACACCCGGCCGAAACTGCT 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               351 CAAT 354
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                                                          ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
                                                                                                                      TOPOLOGY:
                                                                                                                                     STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY
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                                                                                                                                                          nucleic acid
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                                                          Homo sapiens
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Best Loc Matches

171

Query Match

TOPOLOGY:

FILING DATE:

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CAAT 354

746

686 CGAT 683

RESULT 11 US-09-372-498-1/c

SOFTWARE: SEQ ID NO 1

LENGTH: 1342

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US-09-472-880-1/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 120;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 7.3%;
Best Local Similarity 49.2%;
                                                                                                                                                 NAME: Player, William E.
RECISTRATION NUMBER: 31,049
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 1575 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     943 CGAT 940
                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/472,880
FILING DATE: 28-Dec-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Jacobson, Price,
STREET: 400 Seventh Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Daniel CAPUT
Pascale CHALON
Pascual FERRARA
                      FEATURE:
                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Type 2 Neurotensin Receptor (hNT-R2)
                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAAT 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGGCCGCAGCATGATCCTCCGAATCTGGTTGGGCATCCCAGCATACGGCCCAATGTCACAA 350
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                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/FR 9723204 FILING DATE: 17-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Washington STATE: D.C.
                                      ORGANISM: Homo sapiens
                                                                        TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches 124; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Strand, Joann
APPLICANT: Hadcock, John R.
APPLICANT: Hadcock, John R.
TITLE OF INVENTION: No. 6420137el cDNA Encoding The Long and Short
TITLE OF INVENTION: Forms of The Human Neurotensin Subtype 2 Receptor
FILE REFERENCE: 0630/0E792
CURRENT APPLICATION NUMBER: US/09/200,090
CURRENT FILING DATE: 1998-11-25
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1
LENGTH: 1233
TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity 48.8%;
Matches 119; Conservative
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                                      351
                                                                             967
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907 CGAT 904
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                                      CAAT 354
                                                                         AGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTT 290
                                                                                                                                                                                                                                   AGGACACGGCGTTGTAGAGAAGAGGAGTCACAGCTGAGCTGACGTAGAAAAGTATGTTGG
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                                                                                                            TGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAA 350
                                                                                                                                                     TCACCATGTAGAAGTAGTAGAAATTGTACAGTGGGTCAGTCCACGCGTCATCAGGTA
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Pred. No. 0.011;
0; Mismatches 125;
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Pred. No. 0.0044;
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NUMBER OF SEQ ID NOS: 13
SOFTWARE: pt_FL_genes Version 2.0
SEQ ID NO 12
LENGTH: 1535
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (1)..(1338)
US-09-668-680-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Tang, Y. Tom
APPLICANT: Liu, Chenghua
APPLICANT: Zhou, Ping
APPLICANT: Asundi, Vinod
APPLICANT: Asundi, Vinod
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Xue, Aidong J.
APPLICANT: Drmanac, Radoje T.
APPLICANT: Drmanac, Radoje T.
APPLICANT: Drmanac, No. 6436703el Nucleic Acids and
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 790CIP2A
CURRENT APPLICATION NUMBER: US/09/668,680
CURRENT FILING DATE: 2000-09-22
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR FILING DATE: 2000-03-31
PRIOR FILING DATE: 2000-03-31
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US-09-668-680-12/c
: Sequence 12, Application US/09668680
: Patent No. 6436703
: GENERAL INFORMATION:
Search completed: November Job time: 39.4202 secs
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Best Local Similarity 48.6%;
Matches 118; Conservative (
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                                                                                                                                                                                                                                                                                        1172 CACAGGGCCTCCTGGAAGGTCTCTCGGAAGCGGCTGGACATGAGGCTATAGAGCACGGGG 1113
                                                                                              1052 AGG 1050
                                                                                                                                                                                         1112 TTGGCCGCCGAGCCCAGGTAGAAGAAGAAGATGCCGGAGATGACGTGCACGTGCTGGAAGGCC 1053
                                                                                                                                                                                                                                                                                                                    137 CACAGCACCTGCACGAACACCCGGCGAAAACTGCTGCGAGGACACCGTGTACAGGAGGGGG 196
                                                                                                                                                                                                                   197 TTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCC 256
                                                                                                                                            257 CGG 259
                                                                                                                                                                                                                                                                                                                                                                                                                                 77 GAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAG 136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 GACTGGCGCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCGGGCGCTGTCGGTGGTG 76
                    7, 2002, 15:58:19
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Sequence:
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Perfect score:
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Maximum DB seq
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US-09-970-966-210
625
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1: gb_ba:
2: gb_htg
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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

## Result No. იი O 0000000 0000000000000 0000 a a C $\begin{array}{c} 1114 \\ 142 \\ 143 \\ 144 \\ 144 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 \\ 145 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## VERSION KEYWORDS SOURCE ORGANISM RESULT 1 BC011449 LOCUS DEFINITION ACCESSION REFERENCE AUTHORS TITLE JOURNAL BC011449 Homo sapiens, o BC011449 BC011449.1 GI Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1. (bases 1 to 826) Submitted (25-JUL-2001) National Institutes of Health, Mammalian Direct Strausberg, R. Homo sapiens. Homo sapiens Submission GI:15277472 826 bp clone IMAGE:4026092, mRNA mRNA linear PRI 22-AUG-2001

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CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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This clone was selected for full length sequencing because it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
CDNA Library Preparation: Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Institute for Systems Biology
http://www.systemsbiology.org
contact: amadan@systemsbiology.org
Anup Madan, Rachel Dickhoff, Jessica Fahey, Stephanie For
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Contact: MGC help desk
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/clone_lib="NIH_MGC_20"
/lab_host="DH10B-R"
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197 c 228 g
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/db_xref="taxon:9606"
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                                                                                       Submitted (09-JAN-2002) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 63108, USA On Aug 9, 2001 this sequence version replaced g1:14488388.
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Submitted (10-SEP-2000) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St.
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Submitted (09-AUG-2001) Genome Sequencing Center, Washington
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Sulston, J. E. and Waterston, R.
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            Contact: sapiens@watson.wustl.edu
Center project name: H_NH0258B17
                                                         Center: Washington University Genome Sequencing Center Center code: WUGSC
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NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.

This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest.

MAPPING INE Mapping inf McPherson, INFORMATION: information for this on, Department of Gene of Genetics, Washington Un ed by Dr. Joh University, St. D. Louis

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NEIGHBORING SEQUENCE INFORMATION:
The clone sequenced to the left is RP11-11.22; the clone sequenced to the right is RP11-159N20, 2000 bp overlap. Actual start of this clone is at base position 1 of RP11-258B17; actual end is at base position 60003 of RP11-159N20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RP11-258B17 contains a single plasmid region from 1230 to 1239. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOURCE INFORMATION:
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       20104. .20145
/rpt_family="(TG)n"
20170. .20305
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19144
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18933
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16810
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16629
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15532
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941. .1253
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/db_xref="taxon:9606"
/chromosome="2"
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                                                  rpt_family-"ERV1"
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Query Match
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32053..3
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29727. .3
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35361. .35676
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34271. .34348
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26656
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                                                        Score 607; DB 9;
Pred. No. 7.5e-129;
0; Mismatches 1;
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                                                                                     Length 129676;
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TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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Direct Submission
Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7
Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7
Kazusa-Kamatari, Kisarazu, Chiba 292-0812, Japan
(E-mail:genomics@hri.co.jp, Tel:81-438-52-3975, Fax:81-438-52-3986)
NEDO human cDNA sequencing project supported by Ministry of
                                                                                                                                                                                  1

Kawakami, B., Sugiyama, A., Takemoto, M., Sugiyama, T., Irie, R., Otsuki, T., Sato, H., Wakamatsu, A., Ishii, S., Yamamoto, J., Isono, Otsuki, T., Saito, K., Nishikawa, T., Kimura, K., Yamashita, H., Kawai-Hio, Y., Saito, K., Nishikawa, T., Kimura, K., Yamashita, H., Matsuo, K., Nakamura, Y., Sekine, M., Kikuchi, H., Kanda, K., Magatsuma, M., Murakawa, K., Kanehori, K., Takahashi-Fujii, A., Wagatsuma, M., Murakawa, K., Kanehori, K., Magahari, K., Masuho, Y., Nagai
                                                                                                                                                                                                                                                                                                                                                                          AK094501.1 GI:21753575 oligo capping; fis (full insert sequence). Homo sapiens alzheimer cortex cDNA to mRNA,
                                                                                                  Isogai,T. and Yamamoto,J.
                                                                                                                                     Unpublished
                                                                                                                                                                         and Isogai,T.
                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae
                                                                                                                                                                                                                                                                                                                                              Homo
                                                                                                                                                                                                                                                                                                                                                              clone:BRALZ2001350.
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                                                                                                                     (bases 1 to 1832)
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                                                                                                                                                      human cDNA sequencing
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                                                                                  TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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/tissue_type="alzheimer
/clone_lib="BRALZ2"
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/db_xref="GI:21753576"
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/db_xref="taxon:9606"
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KEYWORDS SOURCE ACCESSION VERSION

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COMMENT

Economy, Trade and Industry of Japan; cDNA full insert sequencing: Research Association for Biotechnology (RAB); cDNA library construction: Helix Research Institute (HRI) (supported by Japan Key Technology Center etc.); 5'- & 3'-end one pass sequencing: RAB, HRI, and Biotechnology Center, National Institute of Technology and Evaluation; clone selection for full insert sequencing: HRI and

/translation-"MWVLGIAATFCGLFLLPGFALQIQCYQCEEFQLNNDCSSPEFIV NCTVNVQDMCQKEVMEQSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN TPLCNGPRPKKRGSSASALRPGLRTTILFLKLALFSAHC"

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CCTTGCAGAGGACTGGCGCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCGGGCGCT 66
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Patent: EP 1067182-A 203 10-JAN-2001;
Helix Research Institute (JP)
Location/Qualifiers
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Mammalia; Eutheria; Primates;
1 (bases 1 to 1890)
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/db_xref="GI:14272688"
/db_xref="GI:14272688"
/translation="mwtlgiaatfcglflipgfalqiqcyqceefqlnndcsspefiv
NCTVNVQDMCQKEVMEQSAGINYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNQPRPKKRGSSASALRPGLRTTILFLKLASSRHTAELKEMPPPPALFFQPSPPT
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/db_xref="taxon:9606"
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                      TGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG 606
                                            AGCTTCTTTTGCCACAAGCAAGAGAGAATTTAACACTGTTTCAAACCCGGGGGAGTTGGC
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Patent: EP 1067182-A 620 10-JAN-2001;
Helix Research Institute (JP)
Location/Qualifiers
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Pred. No. 8.8e-100;
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Mammalia; Eutheria; Primates; Catarrhini;
1 (bases 1 to 444)
Kroes, R.A., Moskal, J.R. and Yamamoto, H.
Differential gene expression in cancer
Patent: WO 0136685-A 95 25-MAY-2001;
NYXIS NeuroTheraples, Inc. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGG
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                                  Baker, K.P., Desnoyers, L., Gerritsen, M.E., Goddard, A., Goddowski, P.J., Grimaldi, J.C., Gurney, A.L., Smith, V., Watanabe, C.K. and Wood, W.I.
                                                                                                                                                          Ax358762
Sequence 15 from
Ax358762
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                        Secreted and transmembrane polypeptides
                                 Watanabe,C.
                                                                                   Eukaryota;
Mammalia;
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/db_xref="taxon:9606"
113 c 94 g 14:
                                                                                   ; Metazoa; Chordata; Craniata; Vertebrata; Eutheria; Primates; Catarrhini; Hominidae
0193983-A
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99.5%;
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Pred. No. 2.1e-84;
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. WO0136685
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WO0193983
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                                                                                                Euteleostomi;
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                                                 Stephan, J
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KEYWORDS
SOURCE
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Query Match
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Sequence
AX362255
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                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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15 from Patent WO0208288
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/db_xref="taxon:9606"
433 c 435 g 33
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Location/Qualifiers
                                                    /organism="Homo sapiens"
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433 c 435 g 33
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 62.9%;
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  Score 393; DB 6;
Pred. No. 6.9e-80;
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Sequence 1 from Patent WO0181634.
AX319942
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Methods and compositions for the diagnosis and treatment cardiovascular and tumorigenic disease using 4941
Patent: WO 0181634-A 1 01-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                            Millennium Pharmaceuticals, Inc. (US)
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                                                              Similarity 98.
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LIGNSATIRVTQVLQKKGYLQKEVTDHMVSLACSDILVFLIGMPMEFYSIIWNPLTTS
SYTLSCKLETFLFEACSYATLLHVLTLSFERYLALCHPRYKAVSCPCQVKLLIGFVW
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VFOSSIFGAFVVYLLVVLLSVAFMCWNMMQVLMKSQKGSLAGGTRPQLRKSDSSESRT
VFOSSIFGAFVVYLLVVLLSVAFMCWNMMQVLMKSQKGSLAGGTRPQLRKSDSSESRT
ARROTIFELFLIVVTLAVCHWPNQIRRIMAAAKPKHDWTRSYFRAYMILLEFSETFFY
LSSVINPLLYTVSSQOFRRVFVQVLCCRLSLQHANHEKRLRVHAHSTTDSARFVQRPL
LSSVINPLLYTVSSQOFRRVFVQVLCCRLSLQHANHEKRLRVHAHSTTDSARFVQRPL
LSSVINPLLYTVSSQOFRRVFVQVLCCRLSLQHANHEKRLRVHAHSTTDSARFVQRPL
LFASRRQSSARRTEKIFLSTFQSEAEPQSKSQSLSLESLEPNSGAKPANSAAENGFQE
HEV"
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/protein_id="CAD19381.1"
/db_xref="GI:17901490"
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/db_xref="taxon:9606"
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                                                          Score 361.6; DB 6
Pred. No. 1.1e-72;
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1 (bases 1 to 1362)

McKee, K. K., Tan, C.P., Palyha, O.C., Liu, J., Feighner, S.D., Hreniuk, D.L., Smith, R.G., Howard, A.D. and Van der Ploeg, L.H. Cloning and characterization of two human G protein-coupled receptor genes (GPR38 and GPR39) related to the growth hormone secretagogue and neurotensin receptors

Genomics 46 (3), 426-434 (1997)
                                                                                                                                                                                                                                                                                       Direct Submission
Submitted (17-NOV-1997) Biochemistry and
Inc., PO Box 2000, Rahway, NJ 07065, USA
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                  2 (bases 1 to 1362)
McKee, K.K., Tan, C.P.,
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SYTLSCKLHTFLFEACSYATLLHVLTLSFERVIAICHPERVKAVSGPCQVKLLIGFVW
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                                                                                                                   /note="orphan
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Sequence 3
AX319944
AX319944.1
                                                                                                                                                                                                                                                              Galvin, K.A. and Rudolph-Owen, L.A. Methods and compositions for the diagnosis and treatment cardiovascular and tumorigenic disease using 4941 Patent: WO 0181634-A 3 01-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                              Millennium Pharmaceuticals,
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from Patent W00181634.
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SOURCE KEYWORDS VERSION

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CAGGAAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TGGGCAGA 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CATGATCCTCCGAATCTGGGTTGGGCCATCCAGCATACGGCCAATGTCACAACAATCAGCCT
                                                                                                                                       Direct Submission
Submitted (14-JUN-2002) Genome Sec
Parkway, St. Louis, MO 63108, USA
3 (bases 1 to 209885)
McPherson, J. D. and Waterston, R. H.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                              AC124493
AC124493.2
Center: Washington University Genome Sequencing Center code: WUGSC
Web site:http://genome.wustl.edu/gsc/index.shtml
Contact: submissions@watson.wustl.edu
                                                                                           Submitted (05-JUL-2002) Genome Sequencing Parkway, St. Louis, MO 63108, USA On Jul 5, 2002 this sequence version repla
                                                                                                                                                                                                                                                                                                                                                                                                                                          AC124493 209885 bp DNA linear Mus musculus chromosome UNK clone RP23-462P13, WO SEQUENCE, 5 unordered pieces.
                                                                                                                                                                                                                                   2 (bases 1 to 209885)
McPherson, J.D. and Wat
                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
                                                                                                                                                                                                                                                                    Unpublished
                                                                                                                                                                                                                                                                                    The sequence of Mus musculus
                                                                                                                                                                                                                                                                                                      McPherson, J.D. and
                                                                                                                                                                                                                                                                                                                                     Mammalia;
                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                HTG; HTGS_PHASE1;
                                                                                                                                                                                                                                                                                                                                                                                 house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                   (bases 1 to 209885)
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LSSVINPLLYTVSSQQFRRVFVQVLCCRLSLQHANHEKRLRVHAHSTTDSARFVQRPL
LFASRRQSSARRTEKIFLSTFQSEAEPQSKSQSLSLESLEPNSGAKPANSAAENGFQE
                                                                                                                                                                                                                                                                                                                                   Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                GI:21699722
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          766
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98.9%;
                                                                                                                                                                                                                                       and Waterston, R.H.
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                                                               Genome Center ----
                                                                                                                                                                                                                                                                                                                                                                                                HTGS_DRAFT; HTGS_FULLTOP
                                                                                                                                                                                                                                                                                    Waterston, R.H. s musculus clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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Pred.
                                                                                                                                                                                                         Genome Sequencing
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No. 1.2e-72;
                                                                                                                                                                                                                                                                                                                                  Craniata; Vertebrata;
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          567
                                               Sequencing Center
                                                                                            replaced
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                                                                                                                                                                                                                                                                                                                                     Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          others
                                                                                            gi:21426614
                                                                                                                                                                                                                                                                                                                                                                                                                                                            WORKING DRAFT
                                                                                                                                                                                                         4444 Forest
                                                                                                                             4444 Forest
                                                                                                                                                                                                                                                                                                                                                  Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HTG 05-JUL-2002
                                                                                                                                                                                                                                                                                                                                   Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                         Park
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BASE COUNT
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 451;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
65338 CCACAGAGCTGAGGTAGAAGAAGGTATCAGAGAAGGGCAGGAGGATCATGTATGCCCTGA 65279
                                                                                                                                                                                                           65458 TGAAGCGGGCACGCTGGCGTTTCTCTTGGTTGGCATGCTGCAGAGTCAGGCGGCAGCAGA 65399
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                                                                                                                                                      141 GCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGA 200
                                                                                                                                                                                                                                                                        81
                                                                                                                                                                                                                                                                                                                                                                    21 GGCGCCGGGACGCGAAGAGCAACGGGCGCGCTGCCACAAAGCGGGCGCTGTCGGTGGAGT 80
                                                  TGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGA 260
                                                                                                                                                                                                                                                                                                                  GCCGGGAGGCTAGGAAGATGAGGGGGGCTGCGGGCTGAGCTGGTGCTGTCCTTGGTGGAGA 65459
                                                                                                       GCACCTGCCAGAACACCTTCCGGAACTGCTGAGAGGACACGTTGTAGAGGAGAGGGTTGA
                                                                                                                                                                                                                                                                GCGCATGTACGCGCAGGCGCTTCTCGTGGTGGTGGCGTGCTGCAGCGACAGGCGGCAGCACA 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chemistry: Dye-primer ET; 0% of reads chemistry: Dye-terminator Big Dye; 100% of reads Assembly program: Phrap; version 0,99319 Consensus quality: 208484 bases at least Q40 Consensus quality: 208911 bases at least Q30 Consensus quality: 209285 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Quality coverage: 0.00 in Q20 bases; agarose-fp Quality coverage: 11.62 in Q20 bases; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequencing vector: M13; 0%
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Insert size: 206000; agarose-fp
Insert size: 212112; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequencing vector: plasmid; 100%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Center project name: M_BA0462P13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       59789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NOTE: This is a 'working draft' sequence. It currently consists of 5 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11332
11432
28618
28718
49455
49555
107385
107385
                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="assembly_name:Contig10"
49555. .107384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="assembly_name:Contig12"
44784 c 46164 g 58745 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="assembly_name:Contig11"
107485. .209885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="assembly_name:Contig8"
11432. .28617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculus"
/db_xref="taxon:10090"
/chromosome="UNK"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="assembly_name:Contig9"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="RP23-462P13"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11331: contig of 11331 bp in length
11431: gap of unknown length
28617: contig of 17186 bp in length
28717: gap of unknown length
49454: contig of 20737 bp in length
49554: gap of unknown length
107384: contig of 57830 bp in length
107484: gap of unknown length
107484: gap of unknown length
209885: contig of 102401 bp in length.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           48.3%;
76.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .49454
                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches 131; Indels 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 302; DB 2; Length 209885; Pred. No. 9.4e-59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               403 others
                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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RESULT 13
AC112072
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DEFINITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VERSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACCESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       64980 TCAGGGGACTTGGCTATGTT----AAAAAGCGTTAAATGCTTTCGACAGTGTA 64932
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  65040 CTCTGTGCTGCTGCTTTTTTGCCACTGGCAAAGGAGAATTTAATGCTGCTTCAAGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    65159 AGAAAAGGAGACGGG-GAGAAGAACAGCATGAGAACAAAAATAAATAAATAAAAAAACCCAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65278 AGTACGTTCTGGTCCAGTCATGTTTGGGTTTTGCTGCAGCCATGATCCGTCGGATCTGAT 65219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              556 CCGGGGGAGTTGGCTGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTGNA 608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   438 AAAATATTTAGCCCCCTCTGTTCTG-TGCTTACTGGCCAGGAAATGGTACCAATTTTTCAG 496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            321 TGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAG
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (19-FEB-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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Direct Submission
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                                                                                                                                                                                                                                                   arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                             (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).
NOTE: This is a 'working draft' sequence. It currently
consists of 50 contigs. The true order of the pieces
is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                  NOTE: Estimated insert size may differ from sequence length
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Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 32512 bases at least Q40
Consensus quality: 34515 bases at least Q30
Consensus quality: 35930 bases at least Q20
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                                                                                                                                                                                                                                    SOURCE
                                                                                                                                                                                                                                                    KEYWORDS
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                                                                               TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCTCAGGGGACTTGGCTATGTT----AAAAAGCGTTAAATGCTTTCGACAGTGTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TGTTGGACTTGACAGCTTC---TTTTGCCACAAGCAGAGAGAGAATTTAACACTGTTTCAA 553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TGGGCATCCAGCATACGGCCAATGTCACCAATCAGCCCTGGGCAGACACGAGCAGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACTACGACTTGGTCCAGTCATGTTTGGGTTTTGCTGCGGCCATGATCCGTCGGATCTGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCACGGAGCTGAGGTAGAAGAAGGTGTCGGAGAAGGGGAGGAGGATCATGTACGCCTTGA 75673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAACCTGCCAGAAAACCTTCCGGAACTGCTGAGAAGACACGTTGTAGAGGAGAGGGTTGA 75613
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGAAGTAGGCACGCTGCTTTCTCCTGGTTGGCATGCTGCAGAGTCAGCCGGCAGCAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCCGGGAGGCTAGGAAGATGAGCGGGCTTCGGGGCTGAGCGGCTGCTGTTTTTGGTAGAGC 75493
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACCCGGGGGAGTTGGCTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTGNA 608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGCTGTGCTTGACGGCTTCTTTTTTGCCCACCGGCAAAGGAGAATTTAATGCTGCTTCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAAATATTCAGCCCCTTGGTTCTGTTGCTTACTGGGTAAAAGGGTACCAATCTTTCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAAATATTTAGCCCCTCTGTTCTG-TGCTTACTGGCCAGGAAATGGTACCAATTTTTCAG 496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGAGAGACAGAAAAAGAAAAACACAGCATGAGAACACAGTAAATGAATAAAA----CCAT 437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGT 320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGA 260
                                                                                                                                                                                        AB041649
Mus musculus brain cDNA, clone MNCD-UD/1.
Mus musculus Brain cDNA, clone MNCD-UD/1.
AB041649.1 GI:7670499
AB041649.1 Insert sequence).
fis (full insert sequence).
Mus musculus (strain:C57BL) adult female cDNA to mRNA,
Mus musculus (strain:C57BL) mncb clone:MNCb-0671.
                                                                         Hashimoto, K. isolation of full-length cDNA
                                                                                                                                                       Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
     Hashimoto, K.,
                                       Unpublished
                                                         made by oligo-capping
                                                                                                                 Osada, N., Kusuda, J., Tanuma, R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23927
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/db_xref="taxon:10116"
/clone="CH230-286021"
19653 c 20425 g 23828 t
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1. .92874
                       1 to 1797)
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   Osada, N.,
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                                                             method
   Kusuda,J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches 138;
                                                                                                                                                   Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                               Ito,A.,
   and
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                                                                           mouse brain cDNA library
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                                     CCACAGAGCTGAGGTAGAAGAAGGTATCAGAGAAGGGCAGGAGGATCATGTATGCCCTGA 1617
                                                                                                                                                                                                                                                  TGACCGAGCTGAGGTAGAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGA 260
                                                                                                                                                                                                                                                                                                                            GCACCTGCACGAACACCCGCCGAAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGA 200
                                                                                                                                                                                                                                                                                                                                                                                 TGAAGCGGGCACGCTGGCGTTTCTCTTGGTTGGCATGCTGCAGAGTCAGGCGGCAGCAGA 1497
TGGGCATCCAACACGCCAACGTCACCACAATCAGTCCT-GGCAGGCAAGAACAGGAG
                                                                                                             TGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAG
                                                                                                                                                     AGTACGTTCTGGTCCAGTCATGTTTGGGTTTTGCTGCAGCCATGATCCGTCGGATCTGAT
                                                                                                                                                                       AGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGGCCGAGCCATGATCCTCCGAATCTGGT
                                                                                                                                                                                                                                                                                                        GCACCTGCCAGAACACCTTCCGGAACTGCTGAGAGGACTCGTTGTAGAGGAGAGGGGTTGA
                                                                                                                                                                                                                                                                                                                                                                                                                     GCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCACA 140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1st strand cDNA was primed with an oligo(dT) primer [ATGTGGCCTTTTTTTTTTTTTT]; double-stranded cDNA was synthesized using specific 5' and 3' primers and amplified by PCR. The PCR product was digested with Sfil and size selection was performed to exclude fragments <1.5kb.The Sfil-digested PCR product was cloned into distinct Draill sites of pmE185-FL3. XhoI sites just outside the Draill sites can be used to isolate the cDNA insert. Library was constructed by Sugano et al.(University of Tokyo, Institute of Medical Science). Custom primer used for sequencing (5' end primer [CGACCTGCGAGCCAGA]).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lab host
Vector:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (13-APR-2000) Katsuyuki Hashimoto, National Institute of Infectious Diseases, Division of Genetic Resources; 23-1, Toyama 1-chome, Shinjuku-ku, Tokyo 162-8640, Japan (E-mail:khashienih.go.jp, URL:http://www.nih.go.jp/yoken/genebank/, Tel:81-3-5285-1111(ex.2120), Fax:81-3-5285-1181) URL: http://www.nih.go.jp/yoken/genebank/
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//Protein_1d="BAA95101.1"
//db_xref="GI:7670500"
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TPLCNGPRPKKRGSSASAIRPGLLTTLLFFHLALCLAHC"
a 487 c 511 g 408 t
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/dev_stage="adult"
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/clone="MNCb-0671"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /sex="female"
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Gaps

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RESULT 15 AC128363

190503 bp Di Rattus norvegicus clone CH230-22808, \*\*\*, 69 unordered pieces. AC128363

DNA

NA linear HTG 19-JUL-2002 \*\*\* SEQUENCING IN PROGRESS

SOURCE ORGANISM

Rattus norvegicus Rattus norvegicus. HTG; HTGS\_PHASE1 AC128363.1

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

KEYWORDS VERSION ACCESSION DEFINITION

REFERENCE

Rattus

(bases 1 to 190503)

AUTHORS

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MIZDY, D.M., MAGING, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Ara, J.R., Ayele, M., Banks, T., Barbarla, J., Benton, J., Binage, K., Blankebburg, K., Bonnin, D., Bouck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buck, J., Cholor, S., Chocko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Delancy, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., David, R., Delgado, O., Denn, A.L., Ding, Y., Dinh, K.J., Durbin, K.J., David, R., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Edgar, D., Edwards, G., C., Elhaj, C., Escotto, M., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Edwards, C.C., Elhaj, C., Escotto, M., Edwards, C., Hart, M., Hallowy, C., Hollins, B., Falls, F., Howard, S., Hume, J., Jackson, L.E., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Jackson, L.E., Homsi, F., Howard, S., Huhy, S., Hume, J., Jackson, L.E., Homsi, F., Howard, S., Huhy, S., Hume, J., Liu, W., Louiseged, H., Lozado, R., Jackson, J., Kura, J., Kura, J., Liu, W., Louiseged, H., Lozado, R., Jackson, E., Martin, R., Martindale, A., Martinez, E., Massey, E., Mawhiney, E., McLeer, A., Lucier, R., Luna, R., May, J., Perez, L., Peters, J., Pickers, R., Primus, E., Pull, L., Wolles, M., Ren, Y., Retzer, M., Massey, J., Perez, L., Peters, J., Pickers, R., Primus, E., Pull, J., No, Soon, I., Soon, I., Soon, J., Rohe, M., Sisson, I., Soon, R., Mang, G., Marting, R., Marti
                                                                                                                             Submitted (19-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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JOURNAL REFERENCE AUTHORS TITLE

TITLE

JOURNAL

Direct Submission Worley, K.C. Unpublished 2 (bases 1 Direct Submission

(bases 1 to 190503)

COMMENT

Center code: BCM

Center: Baylor College of Medicine

Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu Contact: hgsc-help@bcm.tmc.edu Center Project Information Center project name: GZCA Center clone name: CH230-22808

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arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                     (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).
NOTE: This is a 'working draft' sequence. It currently consists of 69 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                 NOTE: Estimated insert size may differ from sequence length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequencing vector: Plasmid; Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329 Consensus quality: 131119 bases at least Q40 Consensus quality: 136808 bases at least Q30 Consensus quality: 141839 bases at least Q30
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Best Local Similarity
Matches 233; Conserv
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## ALIGNMENTS

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ABT03280 standard; cDNA; 625

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                                                                 14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
                                                                                                                                                                     cytostatic; gene; ss.
                                                                                                                                                                                            Human ovarian carcinoma associated coding sequence SEQ ID NO:
                                                                                                                                                                                                              05-SEP-2002 (first entry)
                                 Xu J,
                                                 (CORI-) CORIXA CORP
                                                                                                   13-NOV-2001; 2001WO-US45395
                                                                                                                   23-MAY-2002.
                                                                                                                                    WO200239885-A2
                                                                                                                                                    Homo sapiens.
                                                                                                                                                                            Human; ovarian cancer; ovarian carcinoma; gene therapy; immunotherapy;
                2002-500186/53.
                               Stolk JA, Algate PA,
                               Fling SP,
                                 Molesh DA;
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Novel ovarian cancer polypeptide and polynucleotide, useful for

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                     Ovarian
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
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                                         28-JUN-2002
                                                                                 ABL40348 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                        (first entry)
cancer therapy; vaccine;
                   01034C EST clone nucleotide sequence.
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constraint may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polypucleotides of the invention are cuseful for detecting a cancer in a patient, for stimulating and/or ceptagore receils specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for constitution and for determining the presence of a cancer in a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their constitution to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour cells. Polypeptides in tumour cells. Polypeptides and polynucleotides of the current complement of a patient in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current constraint carcinoma ol034C EST clone nucleotides of the constraint carcinoma olose the current constraints.
10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                               (XUJJ/)
(STOL/)
(ALGA/)
(FLIN/)
                                                                                                                                                                                                                                                                                                                                      Ovarian tumour polypeptide and prevention and/or treatment of
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Sequence 625 BP; 182 A; 144 C; 182 G; 116 T; 1 other;

ρ B Š DЪ δÃ В Q В δÃ Matches Query Match Best Local 241 181 181 121 121 61 61 ш GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTTGGCCGCAGC CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG CAGCGACAGGCGCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC CAGCGACAGGCGCAGCACCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG AGTTCTCCTTGCAGAGGACTGGCGCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 625; Similarity Conservative 99.8%; 0; Score 624; Pred. No. Mismatches 4 DB .6e-154; 0 Length 0; Gaps 120 60 300 240 180 180 120 0,

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The present invention describes a composition (1) comprising: carriers and immunostimulants; and a polypeptide (II) of a ovarian tumour polypeptide encoded by a polynucleotide (III) having a cDNA sequence (S1) from the 10912 nucleotide sequences as given in ABL77023 to ABL87934, (III) encoding (II) having a sequence (S2), a T cell population of (II), or antigen presenting cells that express (II). (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to (S1) can be used for detecting ovarian cancer in a patient's biological sample preferably serum or ovarian tissue. The method comprises contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to a predetermined cutoff value and thereby detecting ovarian cancer in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing
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(first entry)

ABT03284 standard;

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cytostatic;

Human; ovarian cancer; ovarian

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Human ovarian carcinoma associated coding sequence SEQ ID NO:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence treatment of ovarian cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer
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03-APR-2001;
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           TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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2001US-0825294.
2001US-0970966.
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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                                                   that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynuclectides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their
         ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma 01034C/0591S consensus
                                                                                                                                                                                                                                         Claim la; Page 127-128; 131pp; English.
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Pred. No. 7.4e-154;
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03-APR-2001; 2001US-0825294
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GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA
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03-APR-2001;
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                                          CAGCGACAGGCGGCAGCAGCACC - TGCACGAACACCCGCCGAAACTGCTGCGAGGACA
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 CCATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC
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2001US-0825294.
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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stream of the entire desired gene or gene fragments, and for designing and
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                                                                                           The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer therapy; vaccine;
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ovarian cancer
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                                                                                                                                                                                     GACAGTGTAAAAAAAAAAAAAAAAA 1617
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                                          cancer;
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57887 given
                                                                                                                                standard;
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                                                                                    (first entry)
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                                       cancer therapy; vaccine; gene therapy; tumour;
                                                                                                                                CDNA; 1619
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99.7%;
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Pred. No. 5.
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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
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                                                      GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAG
CCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC
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                                                                                                                                                                                      CAGCGACAGGCGGCAGCACACCACC-TGCACGAACACCCGCCGAAACTGCTGCTGCGAGGACA 179
                                                                                                                                                                                                                                             GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCCGTGGTTGGCGTGCCTG
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) STOLK J A.
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             The invention relates to 2175 novel human ovarian antigens (ABP41054-ABP43228) and to cDNAs encoding them (ABP4131-ABQ56305), and also encompasses polypeptides 90% identical and polynucleotides 95% identical to the sequences of the invention. The invention additionally relates to recombinant vectors and host cells comprising human ovarian antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABQ54231 standard;
                                                                                                                  Claim
                                                                                                                                                    useful in the prevention, treatment
ovarian cancer), immune disorders,
neurological diseases -
                                                                                                                                                               Isolated nucleic acid molecules encoding novel ovarian polypeptides, useful in the prevention, treatment and diagnosis of cancer (e.g. ovarian cancer), immune disorders, cardiovascular disorders and
                                                                                                                                                                                                                                                                                                                                             07-JUN-2000; 2000US-209467P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of ovarian antigen polynucleotides and polypeptides in diagnosing, treating, prognosing or preventing various ovary and/or breast-related disorders. Such conditions include ovarian cancer and breast cancer, and metastatic tumours of ovarian or breast origin, reproductive system disorders (e.g., infertility, disorders of pregnancy, anovulation, polycystic ovary syndrome, ovarian cysts, and dysmenorrhoea), endocrine disorders, infections (e.g., chlamydia, HJV, toxoplasmosis, and toxic disorders, infections (e.g., chlamydia, HJV, toxoplasmosis, and toxic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      specification,
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                                                                           TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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                                                         Similarity
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99.78;
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Pred. No. 2.
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RESULT 11
AAF22400/c
                         CC cerebroprotective; nootcopic; neuroprotective; antibacterial; virucide; CC fungicide; and ophthalmological. The polynucleotides and proteins can be complete to prevent, treat or ameliorate a medical condition in e.g. humans, cc mice, rabbits, goats, horses, cats, dogs, chickens or sheep. They are compared in diagnosing a pathological condition or susceptibility to a compared in diagnosing a pathological condition or susceptibility to a compared in diagnosing a pathological condition or susceptibility to a compared in diagnose e.g. rheumatol arthritis, hyperproliferative compared in the breast or liver, cardiovascular conditions of the breast or liver, cardiovascular conditions and goal and some set of the breast or liver, cardiovascular conditions caused by bacteria, viruses and fungi and ocular considers e.g. corneal infection. The polypeptides can also be used to disorders e.g. corneal infection. The polypeptides can also be used to call wound healing and epithelial cell proliferation, to prevent skin comporting cell culture of primary tissues, to regenerate tissues and in chemical conditions of preservative to increase or decrease storage capabilities. AAF22364 to AAF22372 and AAB63133 represent sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; ophthalmological; gene therapy; autoimmune disease; neoplasm; rheumatoid arthritis; hyperproliferative disorder; cardiac arrest; cardiovascular disorder; cerebrovascular disorder; cerebral isohaemia; angiogenesis; nervous system disorder; Alzheimer's disease; infection; ocular disorder; corneal infection; wound healing; skin aging;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-APR-1999; 99US-0128694.
20-JAN-2000; 2000US-0176931.
                                                                                                                                                                                                                                                                                                                                                                                                                          and polypeptides homologous to them. Human secreted proteins have activities based on the tissues and cells the genes are expressed in Examples of activities include: immunosuppressive; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-APR-2000; 2000WO-US09071.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid molecule encoding a human secreted protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HUMA-) HUMAN GENOME SCI INC (ROSE/) ROSEN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    F22373 to AAF22421 encode the human secreted proteins given in AAB63182. AAB63183 to AAB63231 represent more human secreted
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DB; AAB63161.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 440; 533pp; English.
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                   invention
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           WO200077026-A1
                                                                                   nootropic; neuroprotective; antibacterial; virucide; fungicide; opthalmalogical; vulnerary; autoimmune disease; cardiovascular disorde hyperproliferative disorders; cerebrovascular disorder; wound healing;
                                                                                                          Human; secreted protein; immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide;
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                                                                                                                                                                                                                                         AAF64188 standard;
                                    Homo sapiens
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                                                                        disorder; skin ageing;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                   Matches 606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; opthalmalogical; and vulnerary. The proteins, polynucleotides, agonists and antagonists can be used to treat or detect or diagnose various diseases and disorders including, autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithalial cell proliferation, to prevent skin ageing due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    culture of primary cissues, to regularize the control of the invention are polypeptides can also be used as a food additive or preservative to polypeptides crease storage capabilities. Included in the invention are polynucleotide sequences AAF64167 - AAF64175 and peptide AAB75505 which polynucleotide sequences AAF64167 - AAF64175 and characterisation of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human secreted proteins AAB75506 - AAB75554 are encoded by polynucleotide sequences AAF64176 - AAF64224. The specification includes amino acid sequences AAB75555 - AAB75606 which represent fragments of the human secreted proteins, and protein sequences with which they share homology.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The proteins and polynucleotides, their agonists and antagonists have activities dependent on the tissues and cells in which they are expressed, examples of these activities include, immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid molecules encoding human secreted proteins, preventing, treating or ameliorating a disorder, e.g. Al Parkinson's diseases and cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1956 BP; 430 A; 543 C; 524 G; 456 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim
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                                                                              504
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                                                                                                                                                                         CAGCGACAGGCGGCAGCACCAGCACCTGCACGACCACCCGCCGAAACTGCTGCGAGGACAC
                                                                                                                                                                                                                                                          GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                   GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
                                                                                                                                                                                                                                      GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                             CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                                                                                                                                       CAGCGACAGCGCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
                                                                                                                                                                                                                                                                                                                    AGTTCTTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAAACGGGCGCTGCACAAAGCG
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                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in the isolation, identification and characterisation of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                    97.18;
99.78;
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                                                                                                                                                                                                                                                                                                                                                                                                 1;
                                                                                                                                                                                                                                                                                                                                                                                               Score 606.6; DB 22
Pred. No. 2.8e-149;
1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                 1.
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                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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RESULT 13
AAF93845
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                                                                                                                                                                                                                                                                                              08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
cDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and diagnosis of diseases associated with inappropriate secretory protein/membrane protein expression. The nucleic acids and complementar
                                                                                 which encode human secretory or membrane proteins represented by AAB88317 - AAB88419. Included in the invention are primers AAF93917 - AAF94295 and AAF62232 - AAF62235 which are used to isolate
                                                                                                            This invention relates to nucleic acid sequences AAF93744 - A which encode human secretory or membrane proteins represented
                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; secretory protein; membrane protein; vaccine; gene therapy; rheumatoid arthritis; diabetes; ss.
                                                                                                                                                                          gene
                                                                                                                                                                                     Nucleic acids encoding
                                                                                                                                                                                                                                                Ota T,
                                                                                                                                                                                                                                                                                                                                               07-JUL-2000; 2000EP-0114090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cDNA encoding a
                                                                                                                                                                                                             P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-MAY-2001
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DB; AAB88418.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACAGTGNA 608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Isogai T,
                                                                                                                                                 SEQ ID
                                                                                                                                                                                                                                                                                              ; 99JP-0194179.
; 2000JP-0118775.
; 2000JP-0183766.
                                                                                                                                                                          or as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                      ding secretory proteins/membrane candidate target molecules in dr
                                                                                                                                                                                                                                                Nishikawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cDNA; 1890
                                                                                                                                               609pp + CD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              membrane or secretory protein clone PSEC0181
                                                                                                                                                                                                                                                Ţ,
                                                                                                                                                 ROM; English
                                                                                                                                                                                                                                                Kawai Y,
                                                                                                                                                                                                                                               Sugiyama T,
                                                                                                                                                                           proteins, useful
rug development -
                                                                                                                                                                                                                                               Hayashi
   complementary
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AAF94186/c
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Human; secretory protein; membrane protein; vaccine; rheumatoid arthritis; diabetes; PCR primer; ss.
                                         Primer specific for DNA encoding secretory/membrane
                                                                                                          AAF94186;
                                                                                                                                       AAF94186 standard;
                                                                                                                                                                                                                  1830
                                                                                                                                                                                                                                                                               1770
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequences may also be used as DNA probes in diagnostic assays (e.g. polymerase chain reactions (PCR)) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and activity. The antibodies and antagonists may also be used as therapeutic agents to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of the polypeptides in samples (e.g. by enzyme linked immunosorbant assay (ELISA). Examples of diseases which may be treated include rheumatoid anthritistic and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                1590
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                                                                                                                                                                                                          CCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCA 366
                                                                                                                                                                                                                                                                                        CATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGAT 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CAGGCGGCAGCACCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTA 186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGA 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCTTGCAGAGGACTGGCGCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCGGGCGCT 66
                                                                                                                                                                                                                                                                                                                                                   AATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACC 486
                                                                                                                                                                                                                                                                                                                                                                                             GACACGAGCAGGAGGAGAGACAGAGAAAAGAAAAACACAGCATGAGAACACAGTAAATA 1709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGAT
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                                                                                                                                                                                                                                                                                                                                  AATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACC
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                                                                           (first
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                                                                                                                                       DNA;
                                                                           entry)
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                                                                                                                                       587
                                                                                                                                       ВP
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Pred. No. 3.9e-147;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; 533 G; 410 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 1890;
                                           protein SEQ
            gene therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                CC which encode human secretory or membrane proteins represented by
CC AARB3317 - AARB4419. Included in the invention are primers
CC AARB3317 - AARB4419. Included in the invention are primers
CC AARB3317 - AARB4419. Included in the invention are primers
CC AARB3317 - AARB4419. Included in the invention are primers
CC CDNA sequences of the invention. The invention also includes methods for
CC Sequences, which can be used in reactions. The polynucleotide sequences
CC can be used in gene therapy. The polynucleotide sequences and the
CC diagnosis of diseases associated with inappropriate secretory
CC protein, membrane protein expression. The nucleic acids and complementary
CC sequences may also be used as DNA probes in diagnostic assays
CC (e.g. polymerase chain reactions (PCR)) to detect and quantitate the
CC used to study the expression and function of secretory proteins/membrane
CC used to study the expression and function of secretory proteins/membrane
CC used to study the include acid sequences in samples. They may also be
CC used to study the expression and function of secretory proteins/membrane
CC used to study the expression and function of secretory proteins/membrane
CC used to study the expression and function of antiponism. The polypeptides may be used
CC dientify modulators (agonists and antagonists) of expression and
CC cidentify. The antipodies and antagonists may also be used as therapeutic
CC agents to down regulate expression and activity. The antibodies may also
CC (ELISA). Examples of diseases which may be treated include rheumatoid
CC arthrities and Alabetos
                                                                                                                                                                                                                                                                                                                                                                 Matches 498;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 587 BP; 117 A; 162 C; 140 G; 162 T; 6 other;
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11-JAN-2000; 2000JP-0118775
02-MAY-2000; 2000JP-0183766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to nucleic acid sequences AAF93744 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acids encoding secretory proteins/membrane pro
        330
                                                                                                                                                                                                                                                                                      516
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                                               338
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                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                   TCGTCCAGTCGTGCTTGGGCTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCC
AGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAGGAGAGACA
                                                                                                                    TGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACC
                                                                                                                                    TGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACC
                                                                                                                                                                                                                        CGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGC
                                                                                                                                                                                                                                                                              CGCNCAGGCGCTTCTCGTGGTTGGCGTGTTCAAGCGACAGCGGCAGCC--AGCACNTGCA
                                                                                                                                                                                                 CGACCACCCGCCGAANTTGTTGCGAGGACACCGTGTACAGAAGCGGGTTGATGACCGAGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and diabetes.
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Pred. No. 4.4e-116;
0; Mismatches 17; Indels 2;
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The present invention describes a composition (I) comprising: carriers can dimmunostimulants; and a polypeptide (II) having a cDNA sequence CC polypeptide encoded by a polypuciectide (III) having a cDNA sequence CC (S1) from the 10912 nucleotide sequences as given in ABI/7023 to CC ABL87934, (III) encoding (II) having a sequence (S2), a T cell CC population of (II), or antigen presenting cells that express (II). CC (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to CC (S1) can be used for detecting ovarian cancer in a patient's biological Sample preferably serum or ovarian tissue. The method comprises CC contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to a predetermined cutoff value and thereby detecting ovarian cancer in the CC patient, where the amount of polynucleotide hybridising to (IV) is CC detected preferably by polymerase chain reaction (PCR). (I) comprising CC (III) and/or (II) is useful for stimulating and/or expanding T cells specific for an ovarian tumour protein comprising contacting T cells with (III) or (II), is useful in design and preparation of cribozyme molecules for inhibiting expression of the tumour polypeptides and proteins in tumour cells; and to isolate a full length gene from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABL79397 standard; cDNA; 558 BP
                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID 2375; 489pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human ovarian cancer related cDNA clone SEQ ID NO:2375
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Matches 539;
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533 NTTAACATTGTTTCAANCCGGGGG
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                                                     CCTGGGCAGACACGAGCAGGAANGGAGAGAGAGAAAAAGAAAAACACAGCATGAGAACA
                                                                                                                                                                             CATTGATTCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGC
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                    TTTAACACTGTTTCAAACCCGGGG
                                          AATGGTACC-ATTTTCAGTGTTTGACTTNACAGCTTCTTTTGCCACAAGCAAGAGAG-A
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Pred. No. 5.
                     561
556
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Search completed: November Job time: 135.06 secs 7, 2002, 10:10:19

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Database
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Maximum DB seq length: 2000000000
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Maximum Match 10
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Gapop 10.0 , Gapext 1.0
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625
24791104 seqs, 12571243825 residues
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                                                 3: /cgn2_6/ptodata/1/pna/US07_COMB.seq: *
5: /cgn2_6/ptodata/1/pna/US080_COMB.seq: *
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Listing first 45 summaries
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/cgn2_6/ptodata/1/pna/US06_COMB.seq:*
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11220.326 Million cell updates/sec
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/cgn2_6,
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result

Result No.	Score	Query Match	Length	DB	ID	Description
1	624		625	- ;	PCT-US01-45395-210	Sequence 210, App
N	624	99.8	625	31	US-09-825-294-210	e 210,
ω	624		625	ω	US-09-867-701-10876	1087
4	624		625	36	US-09-970-966-210	
տ	624		625	42	US-10-212-677-210	210,
6	624		1897	Н	PCT-US01-45395-214	214, A
7	624		1897	3	US-09-825-294-214	Sequence 214, App
8	624		1897	36	US-09-970-966-214	Sequence 214, App
9	624		1897	42	US-10-212-677-214	Sequence 214, App
10	616.4		1967	ட	PCT-US02-29964-16	16, Ar
11	613		1619	ب	PCT-US01-45395-205	•
12	613	98.1	1619	ب	PCT-US01-45395-211	
13	613	98.1	1619	28	US-09-713-550-205	`
14	613	98.1	1619	<u>ω</u>	US-09-825-294-205	`
15	613	98.1	1619	31	US-09-825-294-211	Sequence 211, App
16	613	98.1	1619	36	US-09-970-966-205	205,
17	613	98.1	1619	36	•	211,
18	613	98.1	1619	42	US-10-212-677-205	Sequence 205, App
19	613	98.1	1619	42	US-10-212-677-211	•
20	612.8	98.0	1918	17	US-09-397-022-4454	Sequence 4454, Ap
21	612.8	98.0	1918	31	US-09-808-383-4454	Sequence 4454, Ap

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APPLICANT: COTIAN CORPORATION
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Fling, Steven P.
APPLICANT: Hing, Steven P.
APPLICANT: Molesh, David Alan
APPLICANT: Molesh, David Alan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.48401PC
CURRENT APPLICATION UNMBER: PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEO ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 210
LENCTH: 625
TYPE: DNA
ORGANISM: Homo sapiens
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13 US-10-264-049-111
25 US-09-652-121-729
25 US-09-652-128-9375
26 US-09-652-111-3346
27 US-09-652-11085
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Sequence 1111, App
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Sequence 3346, Ap
Sequence 3458, Ap
Sequence 1608, Ap
Sequence 1714, Ap
Sequence 1814, Ap
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ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(625)
OTHER INFORMATION: n = A
US-09-825-294-210
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APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CF
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT APPLICATION NUMBER: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOTTWARE: FastSEQ for Windows Version 3.0
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Best Local S
Matches 625
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 210
LENGTH: 625
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                                                                                                                    CAGCGACAGGCGGCAGCACCAGCACCTGCACGAACACCCGGCGGAAACTGCTGCGAGGACAC
                                              CAGCGACAGGCGGCAGCACCAGCCACCTGCACGAACACCCGGCCGAAACTGCTGCGAGGACAC
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625; Conserv
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ilarity 100.0%;
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                                                                                                                                                                                                                                                                                                                       Score 624; DB 31;
; Pred. No. 5e-116;
0; Mismatches 0;
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RESULT 1
PCT-US01-45395-210
Sequence 210, Application PC/TUS0145395; GENERAL INFORMATION:

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181

121 61 61 ; FEATURE: ; NAME/KEY: misc\_feature ; LOCATION: 607 ; OTHER INFORMATION: n = PCT-US01-45395-210

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Query Match
Best Local Similarity
Matches 625; Conser

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Sequence 10876, Application US/09867701

GENERAL INFORMATION:
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Jones, Robert
ITILE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
ITILE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
ITILE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILLE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT APPLICATION STORM OF THE SECONDATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10876
LENGTH: 625
TYPE: DNA
ORGANISM: Homo Sapiens
FEATURE:
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; LOCATION: 607
; OTHER INFORMATION: n -
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GENERAL INFORMATION:
APPLICANT: Corixa Corporation
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 214
LENGTH: 1897
                                                                                                                                           Matches 624;
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Best Local
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APPLICANT: Algate, Paul A.
APPLICANT: Molesh, David Alan
APPLICANT: Molesh, David Alan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.48401PC
CURRENT APPLICATION NUMBER: PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 215
                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
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RESULT 7
US-09-825-294-214
Sequence 214, Application US/09825294
GENERAL INFORMATION:
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US-09-825-294-214
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CURRENT FILING DATE: 2001-04-03
NUMBER OF SEO ID NOS: 215
SOFTWARE: FAStSEQ for Windows Version 3.0
SEQ ID NO 214
LENCTH: 1897
TYPE: DNA
                                                                                                                            Matches 624; Conservative
                                                                                                                                        Query Match
Best Local
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APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN
FILE REFERENCE: 210121.484C5
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Pred. No. 6.3e-116;
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 214
LENGTH: 1897
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Best Local
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APPLICANT: Fling, Steven P.
APPLICANT: W1, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C6
CURRENT APPLICATION NUMBER: US/09/970,966
CURRENT FILING DATE: 2001-10-02
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                                                                                                                                           Score 624; DB 36;
Pred. No. 6.3e-116;
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LENGTH: 1897
TYPE: S...
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C7
CURRENT APPLICATION NUMBER: US/10/212,677
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Harlocker, Susan L
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Pred. No. 6.3e-116;
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PRIOR FILING DATE: 2001-09-19
PRIOR APPLICATION NUMBER: US 09/488,725
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: US 09/552,317
PRIOR FILING DATE: 2000-04-25
PRIOR APPLICATION NUMBER: PCT/US00/35017
PRIOR APPLICATION NUMBER: PCT/US00/35017
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
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                                                                                                                                                                                                                                                                               APPLICANT: Haley-Vicente, Dana APPLICANT: Drmanac, Radoje T TITLE OF INVENTION: NOVEL Nucleic TITLE OF INVENTION: Polypeptides FILE REFERENCE: 809ACIP PCT
                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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                                                                  APPLICATION NUMBER: PCT/US01/02623 FILING DATE: 2001-01-25 APPLICATION NUMBER: US 09/496,914
                                                                                                                                       APPLICATION NUMBER: PCT/US00/35017 FILING DATE: 2000-12-22
APPLICATION NUMBER: US 09/560,875 FILING DATE: 2000-04-27 APPLICATION NUMBER: PCT/US01/03800 FILING DATE: 2001-02-05
                                                                                                            APPLICATION NUMBER: US FILING DATE: 2000-01-25
                                                     FILING DATE:
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Zhao, Qing A.
Wang, Jian-Rui
xue, Aidong J.
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Weng, Gezhi
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Ghosh, Malabika
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Best Local S
Matches 617
                                                                     Sequence 205, Application PC/TUS0145395 GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 992
SOFTWARE: pt_FL_genes Version 6.0
SEQ ID NO 16
LENGTH: 1967
TYPE: DNA
                                              APPLICANT: Corixa Corporation APPLICANT: Xu, Jiangchun
  APPLICANT:
                       APPLICANT:
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LOCATION: (338)..(763)
-USO2-29964-16
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PRIOR FILING DATE: 2000-02-28
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Stolk, John A.
Algate, Paul A.
Fling, Steven P.
Molesh, David Alan
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Pred. No. 2.2e-114;
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TITLE

OF.

APPLICANT: APPLICANT: APPLICANT:

T: Algate, Paul A.
T: Fling, Steven P.
T: Molesh, David Alan
INVENTION: COMPOSITIONS

AND

METHODS

FOR THE

THERAPY

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; LENGTH: 1619
; TYPE: DNA
; ORGANISM: Homo s
PCT-US01-45395-205
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Sequence 211, Application PC/TUS0145395
GENERAL INFORMATION:
APPLICANT: Corixa Corporation
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
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TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.48401PC
CURRENT APPLICATION NUMBER: PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 205
LENGTH: 1619
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Best Local Similarity
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                                                                                                                                        CAGCGACAGGCGGCAGCACAGCACC-TGCACGAACACCCGGCGAAACTGCTGCGAGGACA 179
                                                                                                                           GACAGTGTAAAAAAAAAAAAAAAAA 1617
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Sequence 205, Application US/09713550

GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVERLE REFERENCE: 210121.484C4
CURRENT APPLICATION UNUMBER: US/09/713,550
CURRENT FILING DATE: 2000-11-14
NUMBER OF SEQ ID NOS: 205
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; ORGANISM: Homo
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SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 211
LENGTH: 1619
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Best Local Similarity
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APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIA
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION UNMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 205
LENGTH: 1619

CANCER

Sequence 205, Application:

US/09825294

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US-09-825-294-205
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APPLICANT: Xu, Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Algate, Paul A.

APPLICANT: Fling, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CAN

FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 211

LENGTH: 1619

TYPE: DNA

ORGANISM: Homo sapiens
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| Cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
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## ALIGNMENTS

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Query Match	; Remaining Frior Application data ; NUMBER OF SEQ ID NOS: 246 ; SEQ ID NO 15 ; LENGTH: 1524 ; TYPE: DNA ; ORGANISM: Homo Sapien US-10-230-437-15	PRIOR FILING DATE: 1997-12-1; PRIOR FILING DATE: 1998-03-20; PRIOR FILING DATE: 1998-03-20; PRIOR APPLICATION NUMBER: 60, PRIOR FILING DATE: 1998-03-20; PRIOR APPLICATION NUMBER: 60, PRIOR FILING DATE: 1998-03-20; PRIOR APPLICATION NUMBER: 60, PRIOR APPLICATION NUMBER: 60, PRIOR APPLICATION NUMBER: 60, PRIOR FILING DATE: 1998-03-20;	CURRENT APPLICATION NUMBER: US/10/230,437 CURRENT FILING DATE: 2002-08-28 PRIOR APPLICATION NUMBER: 10/119,480 PRIOR FILING DATE: 2002-04-09 PRIOR ETLING DATE: 1997-09-17 PRIOR FILING DATE: 1997-09-17 PRIOR APPLICATION NUMBER: 60/06287 PRIOR APPLICATION NUMBER: 60/063549 PRIOR APPLICATION NUMBER: 60/063549 PRIOR FILING DATE: 1997-10-28 PRIOR FILING DATE: 1997-10-28 PRIOR APPLICATION NUMBER: 60/064103 PRIOR APPLICATION NUMBER: 60/06473	RESULT 1  US-10-230-437-15  Sequence 15, Application US/10230437  GENERAL INFORMATION: GENERAL INFORMATION: GENERAL INFORMATION: APPLICANT: Baker, Kevin P. APPLICANT: Desnoyers, Luc APPLICANT: Gerritsen, Mary APPLICANT: Goddard, Audrey APPLICANT: Goddard, Audrey APPLICANT: Goddard, Audrey APPLICANT: Grimaldi, J. Christophe APPLICANT: Grimaldi, J. Christophe APPLICANT: Surin, Valutin L. APPLICANT: Stephan, Jean-Philippe APPLICANT: Watanabe, Colin L. APPLICANT: Watanabe, Colin L. TITLE OF INVENTION: SECRETED AND THE TITLE DEFENGE. BASCONICA
62.9%; Score 393; DB 6;	removed - see	7 7 7 7 7 7 7 7 7 7 7 7 7	ON NUMBER: US/10/230,437 TE: 2002-08-28  NUMBER: 10/119,480 : 2002-04-09 : 2002-04-09 NUMBER: 60/059113 : 1997-09-17 NUMBER: 60/06287 NUMBER: 60/06287 NUMBER: 60/063549 : 1997-10-28 : 1997-10-28 : 1997-10-31 NUMBER: 60/069873	E.  ANSMEMBRANE THE SAME
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Matches

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Best Local Similarity

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RESULT 2
US-10-270-333-38/c
US-10-270-333-38/c
Sequence 38, Application US/10270333

GENERAL INFORMATION:
APPLICANT: CLEVCHAK, Anibal
TITLE OF INVENTION: INCLEIC G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: MUCLEIC ACID MOLECULES ENCODING GPCR PROTEINS, AND USES
TITLE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
FILE REFERENCE: CL000733CON
CURRENT APPLICATION NUMBER: US/10/270,333
CURRENT FILING DATE: 2002-10-15
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; ORGANISM: Drosophila
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SOFTWARE: FastSEQ for
SEQ ID NO 38
LENGTH: 2196
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PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR FILING DATE: 2000-03-23
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                     183 TGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGA 242
                                                                                                            123 GCGACAGGCGGCAGCACCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCG
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                                                                         GGCCCAGACAAGTGAGCAGCAGCCGCCAGAAGCCGCTGCGAAATTTGGAGGACATCAGGT 1769
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TGTAGAGGATCGGGTTCATGGCTGAGTTTAGGTAGAGCATGAAGCGCGAGAAGTACAGCA 1709
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; OTHER INFORMATION: Human US-60-416-172-67
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, Sequence 67, Application US/60416172
; GENERAL INFORMATION:
, APPLICANT: Chang, Hwai Wen
, APPLICANT: Felts, Katherine A.
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; ORGANISM: Drosophila
US-10-270-333-37
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| Sequence 37, Application US/10270333 |
| GENERAL INFORMATION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | SECURETION: | 
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CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR FILING DATE: 2000-03-23
PRIOR FILING DATE: 2000-03-23
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                                                                                                                                                              SEQ ID NO 67
LENGTH: 667
TYPE: DNA
ORGANISM: Gallus gallus
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SEQ ID NO 37
LENGTH: 6472
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                 APPLICANT: Warren, Amy J.
APPLICANT: Cooper, Judith R.
APPLICANT: Cooper, Judith R.
TITLE OF INVENTION: Genes from Human Cancer Cells and Human
FILE REFERENCE: 216019-153
CURRENT APPLICATION NUMBER: US/60/416,172
CURRENT FILING DATE: 2002-10-04
NUMBER OF SEQ ID NOS: 141
SOFTWARE: Patentin version 3.1
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                                                                               NAME/KEY: misc_feature
                                                                                                                           FEATURE:
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GENERAL INFORMATION:
APPLICANT: Chu, Peter
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                                                                                                                                                                                                                                                                                                                     SEQ ID NO 2
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APPLICANT: Rigel Pharmaceuticals, Inc.
TITLE OF INVENTION: EDG: Modulators of Lymphocyte Activation and Migration
FILE REFERENCE: 021044-000310PC
CURRENT APPLICATION NUMBER: PCT/US02/12342
CURRENT FILING DATE: 2002-08-26
                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/284,763
PRIOR FILING DATE: 2001-04-18
PRIOR APPLICATION NUMBER: US 09/971,228
PRIOR FILING DATE: 2001-10-03
NUMBER OF SEQ ID NOS: 16
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                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                      NAME/KEY: CDS
LOCATION: (9)..(1031)
OTHER INFORMATION: mutant #1 human EDG1
                                                                                                                                                                          FEATURE:
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OTHER INFORMATION:
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121 CAGCGACAGGCGCAGCACAGCACCTGCACGAACACCCGGCGGAAACTGCTGCGAGGACAC 180
                                   Local Similarity 63.4
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Pardo, Jorge
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human endothelial differentiation G-protein
coupled receptor (GPCR) 1 (EDG1)
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                                     Mismatches
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; LOCATION: (1)..(1149)
; OTHER IMFORMATION: wild type human EDG1
PCT-US02-12342-1
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SEQ ID NO 1
LENGTH: 1149
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Best Local Similarity
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APPLICANT:
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PRIOR FILING DATE: 2001-10-03
NUMBER OF SEQ ID NOS: 16
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PRIOR APPLICATION NUMBER: US 60/284,763
PRIOR FILING DATE: 2001-04-18
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                APPLICANT: Spencer, Colin
APPLICANT: Zhao, Haoran
APPLICANT: Rigel Pharmaceuticals, Inc.
APPLICANT: Rigel Pharmaceuticals, Inc.
TITLE OF INVENTION: EDG: Modulators of Lymphocyte Activation and Migration
FILE REFERENCE: 021044-000310PC
CURRENT APPLICATION NUMBER: PCT/US02/12342
                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                         APPLICANT: Chu, Peter APPLICANT: Jiang, Yir
                                                                                                                                                                                         APPLICANT:
CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: wild-type human endothelial differentiation OTHER INFORMATION: G-protein coupled receptor (GPCR) 1 (EDG1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1004 CGGGCACTTGCAGCAGGACATGATCCGGATGAAGGCCCGACGCATCTCCTTGTTGGTCAG 945
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                                                                                                                                                                                                                                                                                                                                                                                                                                            996 CGGGCACTTGCAGCAGGACATGATCCGGATGAAGGCCCGACGCATCTCCTTGTTGGTCAG 937
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                                                                                                                                                                                                                                                                                                                                                                 AGTGTAAATGATGGGGTTGGTGCCGGAGTTGAG
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Zhao, Haoran
Rigel Pharmaceuticals, Inc.
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Rigel Pharmaceuticals, Inc.
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Pardo, Jorge
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                                                                                                                             Masuda, Esteban
Pardo, Jorge
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                                                                                                                                                                       Liao, X.
                                                                                                                                                                                       Li, Congfen
                                                                                                                                                                                                           Jiang, Yingping
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  2002-08-26
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US-10-270-333-97/c
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CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/168,677
PRIOR FILING DATE: 1999-12-03
PRIOR FILING DATE: 1999-12-03
PRIOR REPLICATION NUMBER: 60/175,691
PRIOR RELING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR FILING DATE: 2000-03-23
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: LOCATION: (338)..(1375)
: OTHER INFORMATION: mutant #3 human EDG1
PCT-US02-12342-4
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GENERAL INFORMATION:
APPLICANT: CTAVChik, Anibal
TITLE OF INVENTION: ISOLATED G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GPCR PROTEINS, AND USES
TITLE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
FILE REFERENCE: CL000/333CON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Drosophila
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PRIOR FILING DATE: 2001-04-18
PRIOR APPLICATION NUMBER: US 09/971,228
PRIOR FILING DATE: 2001-10-03
NUMBER OF SEQ ID NOS: 16
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                                                                                    1161
                                                                                                                                                         1221 CGAATAGCGGCACAGCAGGACCTTCTTGAACGCCTGCCGGAACACCTTGTTGAAGATGGT 1162
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                   124 CGACAGGCGGCAGCACCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGT 183
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                                                                                                                                                                                                                                           Local 59;
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                                                                           GTAGAAGATGGGGTTCACCATGGAGCTAGCGTAG 1128
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                                                                                                                                                                                                                                                                    Score 38; DB 6; Length 1251; Pred. No. 0.26;
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Pred. No. 0.19;
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                                                                                                                                                                                                                                              35; Indels
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US-10-270-333-115
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                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                             Query Match
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PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR ETILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR FILING DATE: 2000-03-23
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CURRENT APPLICATION NUMBER: US/10/270,333
CURRENT FILING DATE: 2002-10-15
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TITLE OF INVENTION: ISOLATED G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GPCR PROTEINS, AND USES
TITLE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
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TITLE OF INVENTION: ISOLATED G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GPCR PROTEINS, AND USES
TITLE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
ETLE REFERENCE: CL000733CON
                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
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NUMBER OF SEQ ID NOS: 198
SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/168,677
PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/191,638
                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
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TYPE: DNA
ORGANISM: Drosophila
1431 ACACGAGCTGCTGTAGGCCAGAGTCTGGGCGGTGACCTGGATGACTAGCTTGGTGAGGGT 1490
                                                                                         1371 CACCTTGTAGAAGGCCTTCCGGAAATTCTCGGAGAGGAAGGCGTAGAGCAGCGGATTGAT 1430
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                                      142 CACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGAT 201
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                                                                                                                                                                                                                     Local Similarity
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55.9%;
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                                                                                                                                                                                                         Score 37.4; DB 6; Length 5927; Pred, No. 0.79;
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Pred. No. 0.67;
0; Mismatches 3
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TITLE OF INVENTION: ISOLATED G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GPCR PROTITULE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
FILE REFERENCE: CL000733CON
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/168,677
PRIOR APPLICATION NUMBER: 60/168,677
PRIOR FILING DATE: 1999-12-03
PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR APPLICATION NUMBER: 60/19,638
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR FILING DATE: 2000-03-23
NUMBER OF SEQ ID NOS: 198
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 116
LENGTH: 990
                                                                                                         APPLICANT: Varoglu, M
APPLICANT: He, M
APPLICANT: He, M
APPLICANT: He, M
APPLICANT: He, M
APPLICANT: Sheldon, P
TITLE OF INVENTION: Mitomycin biosynthetic gene cluster
FILE REFERENCE: 600.456US1
CURRENT APPLICATION NUMBER: US/10/267,255
CURRENT FILING DATE: 2002-10-09
PRIOR APPLICATION NUMBER: US 09/266,965
PRIOR FILING DATE: 1996-03-12
PRIOR FILING DATE: 1996-08-19
PRIOR FILING DATE: 1996-08-19
PRIOR FILING DATE: 1994-10-06
PRIOR APPLICATION NUMBER: US 08/133,963
PRIOR APPLICATION NUMBER: US 08/133,963
PRIOR FILING DATE: 1993-10-07
NUMBER OF STOTE NOTE: 1.45
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; ORGANISM: Drosophila
US-10-270-333-116
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                       SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 36 LENGTH: 852
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Best Local Similarity 54.5%;
Matches 73; Conservative
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GENERAL INFORMATION:
APPLICANT: Cravchik, Anibal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -10-267-255-36/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Sherman, D
APPLICANT: Mao, Y
APPLICANT: Varoglu, M
                                                                                                    NUMBER OF SEQ ID NOS:
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                928 GATTGATACACGAGCTGCTGTAGGCCAGAGTCTGGGCGGTGACCTGGATGACTAGCTTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  988 AGAAATTCGCCTTGTAGAAGGCCTTCCGGAAATTCTCGGAGAGGAAGGCGTAGAGCAGCG 929
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; ORGANISM: Streptomyces lavendulae
US-10-267-255-76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   us-10-267-255-76
                                                                                                                                                                                                                                                                                                                                                         Query Match 5.8%; Score 36.2; DB 6; Length 53500; Best Local Similarity 47.9%; Pred. No. 4.7; Matches 104; Conservative 0; Mismatches 113; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 145
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 76
LENGTH: 53500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 76, Application US/10267255 GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/267,255
CORRENT FILING DATE: 2002-10-09
PRIOR APPLICATION NUMBER: US 09/266,965
PRIOR FILING DATE: 1999-03-12
PRIOR APPLICATION NUMBER: US 08/624,447
PRIOR FILING DATE: 1996-08-19
PRIOR FILING DATE: 1994-10-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Varoglu, M
APPLICANT: He, M
APPLICANT: Sheldon, P
TITLE OF INVENTION: Mitomycin biosynthetic gene cluster
FILE REFERENCE: 600.456US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 08/133,963 PRIOR FILING DATE: 1993-10-07
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APPLICANT: Mao, Y
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                                                                                                                                                                          7106 GCGAGCACCTGGGCGCGGTCGGGCATGTGGATGATCGATTCGAGGGCGATGACGGCGTCG
7226 GCGAGCCCGGCCTCCTCGGCCAGCGCGTTGGCCCGGA 7262
                                                                                     7166 AAGCTCTCGTCCTCGAAGGGGAGGTCCATCGCGTCGGCCCGCTGGAAGCGCCCCGGTCG
                                                                                                                                                                                                                  158 CGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAG
                                                                                                               218 AAAAACGTCTCCGAGAAGGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAG
                                        278 TCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAA 314
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    Mismatches

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RESULT 14 PCT-US02-32094-2/c

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RESULT 15

US-10-265-872-2/c

US-10-265-872-2/c

Sequence 2, Application US/10265872

Sequence 2, Application US/10265872

GENERAL INFORMATION:

APPLICANT: Williamson, Mark

TITLE OF INVENTION: METHODS OF USING 241, A HUMAN G-PROTEIN

TITLE OF INVENTION: COUPLED RECEPTOR

FILE REFERENCE: MPIO1-233PIRM

CURRENT APPLICATION NUMBER: US/10/265,872

CURRENT FILING DATE: 2002-10-07

PRIOR APPLICATION NUMBER: 66/328,280

PRIOR FILING DATE: 2001-10-09

NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                ; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 2
; LENGTH: 1285
; TYPE: DNA
; ORGANISM: Homo sapien
US-10-265-872-2
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; TYPE: DNA
; ORGANISM: Homo sapien
PCT-US02-32094-2
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Best Local S
Matches 76
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GENERAL INFORMATION:

APPLICANT: Williamson, Mark

TITLE OF INVENTION: METHODS OF USING 241, A HUMAN G-PROTEIN

TITLE OF INVENTION: COUPLED RECEPTOR

FILE REFERENCE: MPI01-233

CURRENT APPLICATION NUMBER: PCT/US02/32094

CURRENT FILING DATE: 2002-10-08

PRIOR APPLICATION NUMBER: 60/328,280

PRIOR APPLICATION NUMBER: 60/328,280

PRIOR TILING DATE: 2001-10-09

NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2
                                                                                                                                                                                                                                         Query Match
Best Local
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                                                                                                                                            1095 GCGGAGGCACAGAACCTTCTGGAAGCTCTGGCGGAAGTTGTCCGAGAGGAAGCCGTAGAG 1036
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975
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                                 250 GTACGCCCGGAAGTAGGACCTCG 272
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Local Similarity 53.1%;
hes 76; Conservative (
                                                                                                                                                                                                                           Local Similarity
les 76; Conserv
                                                                    GGAGGCGGCTCCTGGGGCAGCG 953
                                                                                                           GAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCAT 249
                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                         5.7%; Score 35.8; 53.1%; Pred. No. 1;
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Search completed: November 7, 2002, 17:57:05 Job time :  $38.4612 \ \text{secs}$ 

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AI990500 ws40d07.x
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## ALIGNMENTS

JOURNAL COMMENT REFERENCE AUTHORS TITLE VERSION KEYWORDS SOURCE ORGANISM RESULT 1 BQ006545/c FEATURES ACCESSION DEFINITION source Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 702)
1 (CGAP) http://www.ncbi.nlm.nih.gov/ncicgap.
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index cDNA Library preparation: Dr. M. Bento Soares, University of Iowa CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa DNA Sequencing by: Dr. M. Bento Soares, University of Iowa Clone Distribution: Clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov POLYA-Yes. Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: Dr. Jose Mercuende BQ006545 702 bp mRNA linear EST 26-MAR-UI-H-EI1-aza-n-21-0-UI.s1 NCI\_CGAP\_EI1 Homo sapiens cDNA clone IMAGE:5846228 3', mRNA sequence.
BQ006545 EST BQ006545.1 GI:19731445 human. Location/Qualifiers EST 26-MAR-2002

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/clone="IMAGE:5846228" /clone\_lib="NCI\_CGAP\_EI1" /tissue\_type="Chondrosarcoma"

/dev\_stage="Adult"

/organism="Homo sapiens" /db\_xref="taxon:9606"

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ACAGTGTAAAAAAAAAAAAAAAAAAA

LOCUS

EST 27-FEB-2002

вм669397/с

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121 CAGCGACAGGCGCAGCACACCTGCACGAACACCCGGCCGAAAACTGCTGCGAGGACAC
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                          ACAGTGNAAAAAAAAAAAAAAAAAAAA
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/lab_host="Duty (Life Technologies)"
/note="Organ: Left Pelvis; Vector: pT7T3-Pac (Pharmacia)
with a modified polylinker; Site_1: EcoR I; Site_2: Not I;
NCI_GAP_EII is a normalized cDNA library containing the
following tissue(s): Chondrosarcoma. The library was
constructed according to Bonaldo, Lennon and Soares,
Genome Research, 6:791-806, 1996. First strand cDNA
synthesis was primed with an oligo-dT primer containing a
Not I site. Double stranded cDNA was ligated to an EcoR I
adaptor, digested with Not I, and cloned directionally
into pT7T3-Pac vector. The oligonucleotide used to prime
the synthesis of first-strand cDNA contains a library tag
sequence that is located between the Not I site and the
(dT)18 tall. The sequence tag for this library is
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  625
                                                                                                                           Local Similarity
                                               1 AGTTCTCCTTGCAGAGGACTGGCGCGGGGCGGGGGGGAGGAGGGAACGGGCGCTGCACAAAGCG 60
Email: msoares@blue.weeg.uiowa.edu
Tissue Procurement: Dr. Greege Hageman
CDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BM669397 627 bp mRNA linear EST 27-UI-E-DW1-ahd-c-11-0-UI.sl UI-E-DW1 Homo sapiens cDNA clone UI-E-DW1-ahd-c-11-0-UI 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  University of Iowa
451 Eckstein Medical Research Building Iowa City, IA 52242, USA
Tel: 319 335 8250
Fax: 319 335 9565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 627) Bonaldo,M.F., Lennon,G. and Soares,M.B. Normalization and subtraction: two approaches to facilitate gene
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BM669397.1 GI:18979294
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                                                                                                                                                                                                                               118
                                                                                                   Conservative
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/note="Organ: eye; Vector: pT7T3-Pac (Pharmacla) with a
modified polylinker; Site_1: EcoR I; Site_2: Not I;
UTE-DW1 is a normalized cDNA library containing the
following tissue(s): lens. The library was constructed
according to Bonaldo, Lennon and Soares, Genome Research,
6:791-806, 1996. First strand cDNA synthesis was primed
with an oligo-dT primer containing a Not I site. Double
stranded cDNA was ligated to an EcoR I adaptor, digested
with Not I, and cloned directionally into pT73-Pac
vector. The oligonucleotide used to prime the synthesis of
first-strand cDNA contains a library tag sequence that is
located between the Not I site and the (dT)18 tail. The
sequence tag for this library is CGATTAGCGA. This library
was created for the program, Gene Discovery in the Visual
                                                                                                                                                                                                                    System, supported by National Eye Institute (NEI).
TAG_LIB=UIE.DW1
TAG_TISSUE-buman lens
TAG_SEO-CGATTAGCGA"
180 c 145 g 183 t 1 others
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/clone_lib="UI-E-DW1"
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/db_xref="taxon:9606"
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1. No. 1.7e-102;
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                                                                                     Email: cgapbs-remail.nih.gov
Tissue Procurement: Invitrogen
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscilence Corporation
Clone distribution: MGC clone distribution: MGC clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 696)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                         AGENCOURT_6507108 NIH_MGC_124
5', mRNA sequence.
                                                          http://image.llnl.gov
Plate: LLAM12722 row: h column:
                                                                                                                                                                                Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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                                  quality sequence stop: 641.
Location/Qualifiers
/organism="Homo sapiens"
/db_xref="taxon:9606"
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He IMAGE:5727798
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RESULT 4 AI936826/c

ACCESSION VERSION

GPR39. ; A1936826

;, mRNA sequence.

A1936826 678 bp mrNA linear EST 08-MAR-2000 wp69h10.x1 NCI\_CGAP\_Brn25 Homo sapiens cDNA clone IMAGE:2467075 3' similar to SW:GP39\_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR

KEYWORDS

AI936826.1 EST.

GI:5675696

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ACAGTGTAAAAAANNNNNAA 685
                                                TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480
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/lab_host="DH10B"
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98.4%;
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Pred. No. 1.6e-100;
0; Mismatches 10;
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CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
                                CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
                                                                                             GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: washington University Genome Clone distribution: NCI-GCAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="anaplastic oligodendroglioma"
/lab_host="DH10B"
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/clone_lib="NCI_CGAP_Brn25"
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/db_xref="taxon:9606"
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99.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      found through the I.M.A.G.E. Consortium/LLNL www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 1212 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AI990500 616 bp mRNA linear EST 08-MAR-2000 ws40d07.xl NCI_CGAP_GC6 Homo sapiens cDNA clone IMAGE:2499661 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1997)
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National Cancer Institute, Cancer Genome Ana
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can b
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/note-"Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Plasmid DNA from the normalized library NCI_CGAP_GC4 was prepared, and scircles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1469064-1470983, and 1475592-1476743). Subtraction by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                   /clone="IMAGE:2499661"
/clone_lib="NCI_CGAP_GC6"
/tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
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Score 601.2;

DB 9;

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REFERENCE
AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
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AW590950

hg51e12.x1 NCI_CGAP_GC6 Homo sapiens cDNA clone IMAGE:2949166 3'
similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A.
R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: M. Bento
                                                                                                   Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 649)
                                                                                                                               Homo sapiens
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AW590950.1 GI:7278094
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                                                                                  National Cancer Institute,
                                                                                          NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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606
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found through the I.M.A.G.E. Consortium/LLNL at:
image.llnl.gov/image/html/iresources.shtml
                                                         DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution
                                                               information
                                                                                       Sequencing Center
                                                               can
                                                               be
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Possible reversed clone: similarity on wrong strand primer: -40UP from quality sequence stop: Location/Qualifiers Gibco 457

ρ /note="vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site\_1: Not I; Site\_2: Eco RI; Plasmid DNA from the normalized library NCI\_CGAP\_GC4 was prepared, and scircles were made in vitro. Following HAP purification this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1469064-1470983, and 1475592-1476743). Subtraction by Bento Scares and M. Fatima Bonaldo. " /tissue\_type="pooled germ
/lab\_host="DH10B" /clone\_lib="NCI\_CGAP\_GC6" /db\_xref="taxon:9606" /clone="IMAGE:2949166" /organism="Homo sapiens" cell tumors"

Similarity Conservative 96.2%; 0, Score 601.2; Pred. No. 7.3e 0; Mismatches .3e-99 DB 10; ω --Indels Length 0, Gaps 0

547 GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60 CAGCGACAGGCGCAGCACAGCACCTGCACGAAACCCCGCCGAAACTGCTGCGAGGACAC GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG AGTTCTCCTTGCAGAGGACTGGCGCCCGGTACGCGAAGATCAACGGGCGCTGCACAAAGCG 180 120 548 488

CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 368 240

428

GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC

308 300

CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC 248 360

TGGGCAGACACGAGCAGGAGGAGAGAGAAAAAGAAAAACACAGCATGAGAACACAG 420 188

TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480 TAAATAAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 128

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KEYWORDS
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        AGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 696)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1997)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          the same 5 libraries. The pools consisted of the following libraries and cloneIDs: Soares NDHSF pool 1: 309384-310919, 323208-325895 Soares NDHSP pool 1: 145032-147335, 147720-148103, 148872-149255, 15002-150407, 151176-152327 Soares NDHSP approximately 1: 758280-760583, 772104-774407 Soares NDHPA pool 1: 304776-306311, 320136-322823, 326280-32663 Soares NDHOT pool 1: 723720-726407, 739080-740999 Subtraction by Bento Soares and M. Fatima Bonaldo."
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/clone_lib="Soares_NSF_F8_9W_OT_PA_P_S1"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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99.5%;
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Pred. No. 8.3e-99;
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690 bp mRNA linear EST 03-NOV-1999
xf40h05.xl NCI_CGAP_Brn50 Homo sapiens cDNA clone IMAGE:2620569 3'
ximilar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Burt Feuerstein, M.D., Mark Israel, M.D.
cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               wrk39. ; mRNA sequence.
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                                                                                                                                                                                                                                           quality sequence stop: 402
/db_xref="taxon:9606"
/clone="IMAGE:2620569"
/clone_lib="NCI_CGAP_Brn50"
/clisue_type="medulloblastoma"
/lab_host="DH10B (phage resistant)"
/note="Organ: brain; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from medulloblastoma tumor tissue, and was then primed with a
                                                                                                                                                                               /organism="Homo sapiens"
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Euteleostomi;

on wrong

oligo(dT) primer. Double-stranded cDNA was ligated

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata, Mammalla; Eutheria; Primates; Catarrhini; Hominidae 1 (Dases 1 to 645)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap National Cancer Institute / National Institute of 1
                                                                                                                     BF432379

Mac53g11 x1 NCI_CGAP_Brn23 Homo similar to SW:GP39_HUMAN 043194 GPR39. , mRNA sequence.

BF432379
                                                                    Homo sapiens
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Pred. No. 1.5e-98;
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Bonaldo, Ph.D.
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Contact: Robert Strausberg,
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                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="NCI_CGAP_Brn23"
/tissue_type="glioblastoma
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /db_xref="taxon:9606"
/clone="IMAGE:3406292"
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98.0%;
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Pred. No. 9.9e-96;
0; Mismatches 12
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GCTGCAGCGACAGCGCAGCACCAGCCACCGAACCACCCGCCGAAACTGCTGCGAGG
                                  AGCGGGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGT 492
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National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BE858216 551 bp mRNA linear EST 29-SEP-7919907.xl NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:3306972 similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEE
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BE858216
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bonaldo, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA Library Preparation: M. Bento Soares, Ph.D.,
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                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cgapbs-r@mail.nih.gov
                                                                                                                                                                                                       /db_xref="taxon:9606"
/clone="IMAGE:3306972"
/clone_lib="NCI_CGAP_Brn23"
/tissue_type="glioblastoma (pooled)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'organism="Homo sapiens"
                                                                                                                                87.7%;
99.8%;
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                                                                                                        Score 548.4; DB 12,
Pred. No. 2.5e-89;
Pred. No. 2.5e-89;
                                                                                                                                               DB 12;
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                                                                                                                                                                                                                                                                                             National Cancer Institute / National Institute of Neurola Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index Unpublished (1998)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
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nac80b12.x1 NCI_CGAP_Brn23 Homo
similar to SW:GP39_HUMAN 043194
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                                                                                                                                             CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 540) NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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Mammalia; Eutheria; Primațes; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                             Tissue Procurement: David N. Louis, M.D.,
                                                                                                                            info@image.llnl.gov
                                                                                                                                                                                                                                        cDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                                            primer: -40UP from Gibco
                                                                                         quality sequence stop:
               /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                     Location/Qualifiers
                                                                                           467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mRNA linear EST 22-JAN-2001 sapiens cDNA clone IMAGE:3440591 3' PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                                                                                                                                                 Myrna R. Rosenfeld M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inear EST 22-JAN-2001 clone IMAGE:3440591 3'
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 537)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.National Cancer Institute / National Institute of Ne Disorders and Stroke, Brain Tumor Genome Anatomy Pro
                                                                                                                                                                                                                      BF594242 537 bp mRNA linear EST 12-DEC-2000 7n10403.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:3564316 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                              Homo sapiens
                                                                                                                                                                                       GPR39.;, mRNA sequence.
BF594242
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="NCI_CGAP_Brn23"
/tissue_type="glioblastoma (pooled)"
/lab_nost="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  86.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 538.4;
Pred. No. 1.6e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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Genome Anatomy Project
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TCAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG
                                             CCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACCAACAACCAGCCCTGGGCAGAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can b
found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tissue Procurement: David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1998)
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="NCI_CGAP_Brn23"
/tissue_type="glioblastoma (pooled)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE: 3564316"
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Best Local Sim:
Matches 534;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: msoares@blue.weeg.ulowa.edu
Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
cDNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
Clone Distribution: Researchers may obtain clones from Research
Genetics (www.resgen.com).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BM667957 538 bp mRNA linear EST 27-FEB-2002 UI-E-DW0-agm-j-01-0-UI.Sl UI-E-DW0 Homo sapiens CDNA clone UI-E-DW0-agm-j-01-0-UI 3', mRNA sequence.

BM667957
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Fax: 319 335 9565
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University of Iowa
451 Eckstein Medical Research Building Iow
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97044477
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Bonaldo, M.F., L
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                                                                           Conservative
                                                                                                                                                                                                                       /lab_host="DHIOB (Life Technologies) (TI phage resistant)"
/note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a
modified polylinker; Site_1: EcoR I; Site_2: Not I;
UI-E-DM0 is a cDNA library containing the following
tissue(s): lens. The library was constructed according to
Bonaldo, Lennon and Soares, Genome Research, 6:791-806,
1996. First strand cDNA synthesis was primed with an
oligo-dT primer containing a Not I site. Double stranded
cDNA was ligated to an EcoR I adaptor, digested with Not
I and cloned directionally into pT7T3-Pac vector. The
oligonucleotide used to prime the synthesis of
first-strand cDNA contains a library tag sequence that is
located between the Not I site and the (dT)18 tail. The
sequence tag for the program, Gene Discovery in the Visual
System, supported by National Eye Institute (NEI).

TAG_ILB=UI-E-DW0
                                                                                                                                                                     TAG_TISSUE-human lens
TAG_SEQ-CGATTAGCGA"
144 c 122 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:9606"

Clone="UI = DWO-agm-j-01-0-UI"

/clone_lib="UI == DWO"

/tissue_type="lens"

/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
                                                                                           85.2%;
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                                                                         Score 532.8; DB 13; Pred. No. 1.6e-86; 0; Mismatches 4;
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                                                                                                              DB 13;
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EATURES SOURCE	AUTHORS TITLE JOURNAL MMENT	ESSION SION WORDS RCE RGANISM	ESULT 14 .1018769/c .OCUS DEFINITION	)y 568 GC 	9y 508 AC      	y 448 GC     b 178 GC	388 CA 	328 CC 	y 268 CC 	y 208 GC 	b 478 CA
. Ph.D.  CDNA Library Arrayed by: Greg Lennon, Ph.D.  CDNA Sequencing by: Washington University Genome Sequencing Center  Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:  www-bio.linl.gov/bbrp/image/lmage.html  Insert Length: 1394 Std Error: 0.00  Seq primer: -40ml3 fwd. ET from Amersham  High quality sequence stop: 452.  Location/Qualifiers  1. 537  /organism="Homo sapiens" /db_xref="taxon:9606" /clone="ib="Soares_testis_NHT" /sex="male" /lab_host="DH10B" /note="Vector: pT7T3D-Pac (Pharmacia) with a modified	Ap http:// al Cancer 3ene Index 5ene (199 ished (199 t: Robert cgapbs-re	AIO18769.1 GI:3232567 AIO18769.1 GI:3232567 EST. buman. Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Eutheria; Primates; Catarrhini; Hominidae; Homo.	769 milar t	GCTGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTGNAAAAAAAAAA	ACAGCTTCTTTGCCACAAGCAAGAGAGAATTTAACACTGTTTCAAACCCGGGGGAGTTG 567 	CCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTTGGACTTG 507	CAGAGAAAAGAAAAAACACAGCATGAGAACACAGTAAATGAAATAAAACCATAAAATATTTA 447 	CAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAGGGAG	CTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCAT 327	CTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGA 267 	CACGAACACCCGCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGA 419

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RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                AW150789 561 bp mRNA linear EST 03-NOV-1999 x939c02.x1 NCI_CGAP_Utl Homo sapiens cDNA clone IMAGE:2629922 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR GPR39. ;, mRNA sequence.
                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 561)
                                Unpublished (1997)
Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                EST.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                                                                                                                                                                                                                                                                                Homo sapiens
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DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortum/LLNL at:
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Emmert-Buck, M.D., Ph.D.
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1 160 c 124 g 158 t 2 others
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/clone_lib="NCI_CGAP_Ut1"
/tissue_type="well-differentiated endometrial
adenocarcinoma, 7 pooled tumors"
/lab_host="H10B"
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## SUMMARIES

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AK094501 .	
LOCUS	AK094501 1832 bp mRNA linear PRI 15-JUL-2002
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	to Homo sapiens Gz-selective GTPase-activating protein (RGS20)
	mRNA.
ACCESSION	AK094501
VERSION	AK094501.1 GI:21753575
KEYWORDS	oligo capping; fis (full insert sequence).
SOURCE	Homo sapiens alzheimer cortex cDNA to mRNA, clone_lib:BRALZ2
	clone:BRALZ2001350.
ORGANISM	Homo sapiens
	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE	<b></b>

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CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCAGGCCAGGGCTCCGCAC
                                                                                                                                                     CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
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                                                                                                                                       CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAA
                                                                                                                                                                                                                                                                          GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA
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/codon_start=1
/protein_id="BaC04368.1"
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NCTVNVQDMCQKEVMEDSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRPKKRGSSASALRPGLRTTILFLKLALFSAHC"
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/clone_lib="BRAL22"
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/clone="BRALZ2001350"
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                                 CCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTGGACTTGA
                                                                                 AGAGAAAAGAAAAACACAGCATGAGAACACAGTAAATGAATAAAACCATAAAATATTTAG
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                   CCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTGGACTTGA
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                                                                                                                all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
                                                                                                                                                                                                                                                                                          NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (09-JAN-2002) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 6. On Aug 9, 2001 this sequence version replaced gi:14488388.
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Submitted (09-AUG-2001) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St.
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Homo sapiens BAC clone RP11-258B17
AC079773
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5 (bases 1 to 129676)
Waterston, R.
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Mapping information for this clone was provided by Dr. John D McPherson, Department of Genetics, Washington University, St.
                                                  MAPPING INFORMATION:
                                                                                                                                                                                                                                          This sequence was finished as follows unless otherwise noted:
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                                                                                                   restriction digest.
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Sulston, J.E. and Waterston, R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Web site: http://genome.wustl.edu/gsc
Contact: sapiens@watson.wustl.edu
Contact: Summary Statistics
                                                                                                                                                                                                                                                                                                                                                                                                                                           Center project name: H_NH0258B17
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VECTOR: pBACe3.6
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The clone sequenced to the left is RP11-1L22; the clone sequenced to the right is RP11-159N20, 2000 bp overlap. Actual start of this clone is at base position 1 of RP11-258B17; actual end is at base clone is at base position 1 of RP11-258B17; actual end is at base
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MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc \,
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18434. .18801
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                                                                                  TGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCC 1249
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80.9%; Score 1310; DB 9; Length 129676; ty 99.9%; Pred. No. 0; ervative 0; Mismatches 0; Indels 1; Gaps 1;	.35676 amily="Alu"	6. Marchi de Esi bessosoi (Mid:9143/40/2) 6. 35374 familya"/TTTTC):	camily "MERI_Cype"3470336703	34348	74PC_Lamily="Mix" 34079amily="Mix" 7594 for=11:-#15#	32787	32080	7 A A	/PC_1dHI1Y="MEK1_Cype" 3149431783 /************************************	באז ניייס	מים מים	29727. 30221 29727. 30221	: 5	70000 march to Est Br/300/0 (MID-912002/44) 28834 . 28958 7004 femiliary mares	D EST DE736070 (NTD. 213063744	/FPC_IdHIIY="(TA)H" 2792628052 /************************************	ma"	7 F	7 PC_10HI1Y="L1" 2740327671 //// f527671	1 m / m	/pt_ramily="Mek2_type" 272463 272463	י מולא	ER2	RVL"	/rpt_tamily="MaLR" 2641726463	/rpt_tamily="MIR" 2548525805	/rpt_family="Alu" 2510825208	7:pe_ramily="Mik" 2396724256	//OCTE="MATCH TO EST BF821//9 (NID:912151303)" 23084. 23228	camily="(A)n" 21796	Tam11y="MER1	_family="MER1_t 820756

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GGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGG

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Primates; Craniata; Vertebrata; Catarrhini; Hominidae; DNA and Euteleostomi; 14-DEC-2001 1459 1320 1279 1140 1220 1020 1260 1339 1200 1080 1160 1040 840 920 780 860 720